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- (54) Glycopeptide antibiotic derivatives.
- 57) The present invention provides glycopeptide antibiotic derivative compounds. These derivative compounds possess antibacterial activity against a wide variety of bacteria, including activity against vancomycin-resistant isolates. Methods of making and using these glycopeptide antibiotic derivative compounds are also provided.

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N w improv d antibiotics ar continually in d mand, particularly for the treatment of human dis as s. Increased potency, expanded spectrum f bacterial inhibition, increased in vivo efficacy, and improved pharmaceutical properties are som of thing also for improved antibiotics.

In the s arch for n w antibiotics, structural modification of known antibi tics is attempted when ver possible. The glycopeptide antibiotics have such complex structures that even small changes are difficult. Furthermore, it is difficult to predict the effect these changes will make in the antimicrobial and physiological properties. Processes for modifying known antibiotics and the new active derivatives made by such processes, therefore, continue to be of great importance.

Previously, N-alkyl and N-acyl derivatives of the glycopeptides vancomycin, A51568A, A51568B, M43A and M43D have been prepared (U.S. Patent Nos. 4,639,433, 4,643,987, and 4,698,327). Several of these compounds exhibited microbiological activity, including activity against vancomycin-resistant isolates. Nicas et al., Antimicrobial Agents and Chemotherapy, 33(9):1477-1481 (1989). In addition, European Patent Application Publication No. 0435503, published July 3, 1993, describes certain N-alkyl and N-acyl derivatives of the A82846 glycopeptides, factors A, B, and C.

The formula I compounds of this invention are new members of the glycopeptide group of antibiotics. These new compounds are derivatives of known glycopeptide antibiotics that include vancomycin (U.S. Patent 3,067,099); A82846A, A82846B, and A82846C (U.S. Patent 5,312,738, European Patent Publication 256,071 AI); PA-42867 factors A, C, and D (U.S. Patent 4,946,941 and European Patent Publication 231,111 A2); A83850 (U.S. Patent No. 5,187,082); avoparcin (U.S. Patent 3,338,786 and U.S. Patent 4,322,343); actinoidin, also known as K288 (J. Antibiotics Series A 14:141 (1961); helevecardin (Chem. Abstracts 110:17188 (1989) and Japanese Patent Application 86/157,397); galacardin (Chem. Abstracts 110:17188 (1989) and Japanese Patent Application 89/221,320); and M47767 (European Patent Publication 339,982). The references listed above which describe these glycopeptides are incorporated herein by reference.

Enterococci are important human pathogens. Infections caused by enterococci are generally difficult to treat. Glycopeptides, such as vancomycin and teicoplanin, have become important therapies in the treatment of infections due to enterococci. However, strains of Enterococcus faecium and E. faecalis have recently been isolated that are resistant to vancomycin and teicoplanin. Leclercq et al., "Plasmid Mediated Resistance to Vancomycin and Teicoplanin in Enterococcus Faecium," <a href="The New England Journal of Medicine, 319(3):157-161 (1988), and Uttley et al., "Vancomycin-Resistant Enterococci," Lancet, 1:57-58 (1988). The isolates were also found to be resistant to other antibiotics. A recent survey found 7.9% of Enterococci in United States hospitals are now vancomycin resistant. "Nosocomial Enterococci Resistant to Vancomycin" Morbidity and Mortality Weekly Report 42 (30):597-598 (1993). In addition to their broad activity against gram-positive organisms, many of the glycopeptide compounds of this invention also exhibit improved antimicrobial activity against vancomycin-resistant isolates.

The present invention provides compounds of the formula I:

or salt thereof, wherein:

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X and Y are ach indep indently hydrogen or chloro:

R is hydrogen, 4- pi-vancosaminyl, actin saminyl, or ristosaminyl;

R1 is hydrogen, r mannose;

 R^2 is $-NH_2$, $-NHCH_3$, or $-N(CH_3)_2$;

 R^3 is $-CH_2CH(CH_3)_2$, [p-OH, m-Cl]ph nyl, p-rhamnos -ph nyl, r[p-rhamnose-galactos]ph nyl, [p-ga-5 lactose-galactose]phenyl, [p-CH₃O-rhamnose]phenyl;

R4 is -CH₂(CO)NH₂, benzyl, [p-OH]phenyl, or [p-OH, m-Cl]phenyl;

R⁵ is hydrogen, or mannose;

R6 is 4-epi-vancosaminyl, L-acosaminyl, L-ristosaminyl, or L-actinosaminyl;

R7 is (C2-C16)alkenyl, (C2-C12)alkynyl, (C1-C12 alkyl)-R8, (C1-C12 alkyl)-halo, (C2-C6 alkenyl)-R8, (C2-C6 alkynyl)-R₈, (C₁-C₁₂ alkyl)-O-R₈, and is attached to the amino group of R⁶;

R8 is selected from the group consisting of:

- a) multicyclic aryl unsubstituted or substituted with one or more substituents independently selected from the group consisting of:
 - (i) hydroxy,
 - (ii) halo,

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- (iii) nitro,
- (iv) (C₁-C₆)alkyl,
- (v) (C₁-C₆)alkenyl,
- 20 (vi) (C1-C6)alkynyl,
 - (vii) (C1-C6)alkoxy,
 - (viii) halo-(C1-C6)alkyl,
 - (ix) halo-(C1-C6)alkoxy,
 - (x) carbo-(C₁-C₆)alkoxy,

 - (xi) carbobenzyloxy,
 - (xii) carbobenzyloxy substituted with (C₁-C₆)alkyl, (C₁-C₆)alkoxy, halo, or nitro,
 - (xiii) a group of the formula -S(O)n-R9, wherein n' is 0-2 and R9 is (C1-C6)alkyl, phenyl, or phenyl substituted with (C1-C6)alkyl, (C1-C6)alkoxy, halo, or nitro, and
 - (xiv) a group of the formula -C(O)N(R10)2 wherein each R10 substituent is independently hydrogen, (C1-
 - C₆)-alkyl, (C₁-C₆)-alkoxy, phenyl, or phenyl substituted with (C₁-C₆)-alkyl, (C₁-C₆)-alkoxy, halo, or nitro;
 - b) heteroaryl unsubstituted or substituted with one or more substituents independently selected from the group consisting of:
 - (i) halo,
 - (ii) (C1-C6)alkyl,
 - (iii) (C1-C8)alkoxy,
 - (iv) halo-(C1-C6)alkyl,
 - (v) halo-(C₁-C₆)alkoxy,
 - (vi) phenyl,
 - (vii) thiophenyl,
 - (viii) phenyl substituted with halo, (C₁-C₆)alkyl, (C₁-C₆)alkenyl, (C₁-C₆)alkynyl, (C₁-C₆)alkoxy, or nitro,
 - (ix) carbo-(C₁-C₆)alkoxy,
 - (x) carbobenzyloxy,
 - (xi) carbobenzyloxy substituted with (C₁-C₆)alkyl, (C₁-C₆) alkoxy, halo, or nitro,
 - (xii) a group of the formula -S(O),-R9, as defined above,
 - (xiii) a group of the formula -C(O)N(R10)2 as defined above, and
 - (xiv) thienyl;
 - c) a group of the formula:

wherein A^1 is $-OC(A^2)_2-C(A^2)_2-O$, $-O-C(A^2)_2-O$, $-C(A^2)_2-O$, or $-C(A^2)_2-C(A^2)_2-C(A^2)_2-C(A^2)_2$, and 55 ach A2 substituent is independently s I ct d from hydrogen, (C1-C6)-alkyl, (C1-C6)alk xy, and (C4-C10)cycloalkyl;

d) a group of the formula:

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wherein p is from 1 to 5; and

R¹¹ is independently selected from the group consisting of:

- (i) hydrogen,
- (ii) nitro,
- (iii) hydroxy,
 - (iv) halo,
 - (v) (C₁-C₈)alkyl,
 - (vi) (C1-C8)alkoxy,
 - (vii) (C₉-C₁₂)alkyl,
 - (viii) (C2-C9)alkynyl,
 - (ix) (C₉-C₁₂)alkoxy,

 - (x) (C₁-C₃)alkoxy substituted with (C₁-C₃)alkoxy, hydroxy, halo(C₁-C₃)alkoxy, or (C₁-C₄)alkylthio,
 - (xi) (C₂-C₅)alkenyloxy,
 - (xii) (C₁-C₁₃)alkynyloxy
- (xiii) halo-(C1-C6)alkyl,
 - (xiv) halo-(C₁-C₆)alkoxy,
 - (xv) (C2-C8)alkylthio,
 - (xvi) (C2-C10)alkanoyloxy,
 - (xvii) carboxy-(C2-C4)alkenyl,
 - (xviii) (C1-C3)alkylsulfonyloxy,

 - (xix) carboxy-(C₁-C₃)alkyl,
 - (xx) N-[di(C_1 - C_3)-alkyl]amino-(C_1 - C_3)alkoxy,
 - (xxi) cyano-(C1-C6)alkoxy, and
 - (xxii) diphenyl-(C1-C6)alkyl,

with the proviso that when R11 is (C1-C8)alkyl, (C1-C8)alkoxy, or halo, p must be greater or equal to 2, or when R7 is (C1-C3 alkyl)-R8 then R11 is not hydrogen, (C1-C8)alkyl, (C1-C8)alkoxy, or halo;

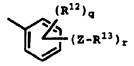
e) a group of the formula:

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40 wherein q is 0 to 4;

R¹² is independently selected from the group consisting of:

- (i) halo,
- (ii) nitro,
- (iii) (C1-C8)alkyl,
- (iv) (C₁-C₆)alkoxy,
- (v) halo-(C₁-C₆)alkyl,
- (vi) halo-(C₁-C₆)alkoxy, and
- (vii) hydroxy, and
- (vii) (C₁-C₆)thioalkyl;

r is 1 to 5; provided that the sum of q and r is no greater than 5;

Z is selected from the group consisting of:

- (i) a single bond,
- (ii) divalent (C₁-C₆)alkyl unsubstituted or substituted with hydroxy, (C₁-C₆)alkyl, or (C₁-C₆)alkoxy,
- (iii) divalent (C2-C6)alkenyl,
- (iv) dival nt (C2-C6)alkynyl, or

(v) a group of th formula -(C(R14)₂)_s-R15- or -R15-(C(R14)₂)_s-, wherein s is 0-6; wher in ach R14 substitu nt is ind pendently s lected from hydrogen, (C₁-C₆)-alkyl, or (C₄-C₁₀) cycloalkyl; and R¹⁵ is sel ct d from -O-, -S-, -SO-, -SO₂-, -SO₂-O-, -C(O)-, -OC(O)-, -C(O)O-, -NH-, -N(C₁-C₆ alkyl)-, and -C(O)NH-, -NHC(O)-, N=N;

R13 is independ ntly selected from the grup consisting of:

- (i) (C₄-C₁₀)heterocyclyl,
- (ii) heteroaryl,
- (iii) (C₄-C₁₀)cycloalkyl unsubstitut d or substituted with (C₁-C₆)alkyl, r
- (iv) phenyl unsubstituted or substituted with 1 to 5 substituents independently selected from: halo, hydroxy, nitro, (C_1-C_{10}) alkyl, (C_1-C_{10}) alkoxy, halo- (C_1-C_3) alkoxy, halo- (C_1-C_3) alkyl, (C_1-C_3) alkoxyphenyl, phenyl- (C_1-C_3) alkyl, (C_1-C_6) alkyl, (C_1-C_6) alkyl, phenyl- (C_1-C_6) alkyl, (C_1-C_6) alkylphenyl;
- f) (C₄-C₁₀)cycloalkyl unsubstituted or substituted with one or more substituents independently selected from the group consisting of:
 - (i) (C₁-C₆)alkyl,
 - (ii) (C₁-C₆)alkoxy,
 - (iii) (C1-C6)alkenyl,
 - (iv) (C₁-C₆)alkynyl,
 - (v) (C₄-C₁₀)cycloalkyl,
 - (vi) phenyl,
 - (vii) phenylthio,
 - (viii) phenyl substituted by nitro, halo, (C1-C6)alkanoyloxy, or carbocycloalkoxy, and
 - (ix) a group represented by the formula -Z-R13 wherein Z and R13 are as defined above; and
- g) a group of the formula:

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wherein

A3 and A4 are each independently selected from

- (i) a bond,
- (ii) -O-,
- (iii) -S(O),-, wherein t is 0 to 2,
- (iv) $-C(R^{17})_2$, wherein each R^{17} substituent is independently selected from hydrogen, (C_1-C_6) alkyl, hydroxy, (C_1-C_6) alkyl, (C_1-C_6) alkoxy, or both R^{17} substituents taken together are O,
- (v) -N(R¹⁸)₂-, wherein each R¹⁸ substituent is independently selected from hydrogen; (C₁-C₆)alkyl; (C₁-C₆)alkenyl; (C₁-C₆)alkynyl; (C₄-C₁₀)cycloalkyl; phenyl; phenyl substituted by nitro, halo, (C₁-C₆)alkanoyloxy; or both R¹⁸ substituents taken together are (C₄-C₁₀)cycloalkyl;

R¹⁶ is R¹² or R¹³ as defined above; and

u is 0-4.

Another aspect of the invention relates to compositions for the treatment of susceptible bacterial infections comprising a compound of formula \underline{I} in combination with an acceptable pharmaceutical carrier. Methods for the treatment of susceptible bacterial infections with compositions of formula \underline{I} are also a part of this invention.

The alkyl substituents recited herein denote substituted or unsubstituted, straight or branched chain hydrocarbons of the length specified. The term "alkenyl" refers to a substituted or unsubstituted, straight or branched alkenyl chain of the length specified. The term "alkynyl" refers to a substituted or unsubstituted, straight or branched alkynyl chain of the length specified.

The alkoxy substituents recited herein represent an alkyl group attached through an oxygen bridge. The term "alkenoxy" represents a alkenyl chain of the specified length attached to an oxygen atom.

The term "multicyclic aryl" means a stable, saturated or unsaturated, substituted or unsubstituted, 9 to 10 membered organic fused bicyclic ring; a stable, saturated or unsaturated, substituted or unsubstituted 12 to 14 membered organic fused tricyclic ring; or a stable, saturated or unsaturated, substituted or unsubstituted 14 to 16 membered organic fused tetracyclic ring. The bicyclic ring may have 0 to 4 substituents, the tricyclic ring may have 0 to 6 substituents, and the tetracyclic ring may have 0 to 8 substituents. Typical multi-cyclic aryls include fluorenyl, napthyl, anthranyl, phenanthranyl, biphenylene and pyrenyl.

The term "heteroaryl" represents a stable, saturated or unsaturated, substituted or unsubstituted, 4 to 7 membered rganic m nocyclic ring having a het r atom selected from S, O, and N; a stabl., saturated or unsaturat d, substituted or unsubstitut d, 9 to 10 m mbered organic fused bicyclic ring having 1 to 2 h tero atoms s. I cted from S, O, and N; or a stable, saturated or unsaturat d, substituted or unsubstituted, 12 to 14 membered organic fus. d tricyclic ring having a h. t. ro atom selected from S, O, and N. The nitrog. n and sulfur

atoms of thes rings are optionally oxidiz d, and the nitrog n h tero atoms ar opti nally quarternized. Th monocyclic ring may have 0 to 5 substitu nts. The bicyclic ring may have 0 to 7 substitu nts, and the tricyclic ring may have 0 to 9 substituents. Typical heteroaryls include quinolyl, piperidyl, thienyl, pip ronyl, xaflu renyl, pyridyl and b nzothienyl and th lik.

The term " (C_4-C_{10}) cycloalkyl" mbraces substitu nts having from four to ten carbon atoms, such as cyclobutyl, cyclopentyl, cyclohexyl, and cycloheptyl which may be unsubstituted or substituted with substituents such as alkyl and phenyl. This term also embraces C_5 to C_{10} cycloalkenyl groups such as cyclopentenyl and cyclohexenyl. The term " (C_4-C_{10}) cycloalkyl" also embraces bicyclic and tricyclic cycloalkyls such as bicyclopentyl, bicyclohexyl, bicycloheptyl, and adamantyl.

The term "alkanoyloxy" represents an alkanoyl group attached through an oxygen bridge. These substituents may be substituted or unsubstituted, straight, or branched chains of the specified length.

The term "cyano-(C₁-C₆)alkoxy" represents a substituted or unsubstituted, straight or branched alkoxy chain having from one to six carbon atoms with a cyano moiety attached to it.

The term "divalent (C_1 - C_6)alkyl" represents an unsubstituted or substituted, straight or branched divalent alkyl chain having from one to six carbon atoms. Typical divalent (C_1 - C_6)alkyl groups include methylene, ethylene, propylene, isopropylene, butylene, isobutylene, sec-butylene, t-butylene, pentylene, neo-pentylene, and hexylene. Such divalent (C_1 - C_6)alkyl groups may be substituted with substituents such as alkyl, alkoxy, and hydroxy.

The term "divalent (C₂-C₆)alkenyl" represents a straight or branched divalent alkenyl chain having from two to six carbon atoms. Typical divalent (C₂-C₆)alkenyl include ethenyl, 1-propenyl, 2-propenyl, 1-butenyl, 2-butenyl and the like.

The term "divalent (C_2 - C_6)alkynyl" represents a straight or branched divalent alkynyl chain having from two to six carbon atoms. Typical divalent (C_2 - C_6)alkynyl include ethynylene, 1-propynylene, 2-propynylene, 1-butynylene, 2-butynylene and the like.

The term "halo" represents chloro, fluoro, bromo or iodo.

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The term "halo-(C₁-C₆)alkyl" represents a straight or branched alkyl chain having from one to six carbon atoms with from 0 to 3 halogen atoms attached to each carbon. Typical halo-(C₁-C₆)alkyl groups include chloromethyl, 2-bromoethyl, 1-chloroisopropyl, 3-fluoropropyl, 2,3-dibromobutyl, 3-chloroisobutyl, iodo-t-butyl, tri-fluoromethyl, and the like.

The term "halo- (C_1-C_6) alkoxy" represents a straight or branched alkoxy chain having from one to six carbon atoms with from 0 to 3 halogen atoms attached to each carbon. Typical halo- (C_1-C_6) alkoxy groups include chloromethoxy, 2-bromoethoxy, 1-chloroisopropoxy, 3-fluoropropoxy, 2,3-dibromobutoxy, 3-chloroisobutoxy, iodo-t-butoxy, trifluoromethoxy, and the like.

The term "heterocycly!" embraces saturated groups having three to ten ring members and which heterocyclic ring contains a hetero atom selected from oxygen, sulfur and nitrogen, examples of which are piperazinyl, morpholino, piperdyl, methylpiperdyl, azetidinyl, and aziridinyl.

The invention includes salts of the compounds defined by formula I. Although generally neutral, a compound of this invention can possess a sufficiently acidic, a sufficiently basic, or both functional groups, and accordingly react with any of a number of inorganic bases, and inorganic and organic acids, to form a pharmaceutically acceptable salt.

The term "pharmaceutically acceptable salt" as used herein, refers to salts of the compounds of the above formula I which are substantially non-toxic to living organisms. Typical pharmaceutically acceptable salts include those salts prepared by reaction of the compounds of the present invention with a pharmaceutically acceptable mineral or organic acid or an inorganic base. Such salts are known as acid addition and base addition salts.

Acids commonly employed to form acid addition salts are inorganic acids such as hydrochloric acid, hydrobromic acid, hydroiodic acid, sulfuric acid, phosphoric acid, and the like, and organic acids such as p-toluenesulfonic acid, methanesulfonic acid, oxalic acid, p-bromophenylsulfonic acid, carbonic acid, succinic acid, citric acid, benzoic acid, acetic acid, and the like. Examples of such pharmaceutically acceptable salts are the sulfate, pyrosulfate, bisulfate, sulfite, bisulfite, phosphate, monohydrogenphosphate, dihydrogenphosphate, metaphosphate, pyrophosphate, chloride, bromide, iodide, acetate, propionate, decanoate, caprylate, acrylate, formate, isobutyrate, caproate, heptanoate, propiolate, oxalate, malonate, succinate, suberate, sebacate, fumarate, maleate, butyne-1,4-dioate, hexyne-1,6-dioate, benzoate, chlorobenzoate, methylbenzoate, dinitrobenzoate, hydroxybenzoate, methoxybenzoate, phthalate, sulfonate, xylenesulfonate, phenylacetate, phenylpropionate, phenylbutyrate, citrat, lactate, g-hydroxybutyrat, glycollate, tartrate, methanesulfonate, propanesulfonate, naphthalene-1-sulf nate, naphthal ne-2-sulfonate, mandelate and the like. Preferr d pharmaceutically acceptable acid addition salts are those formed with mineral acids such as hydrochloric acid and hydrobromic acid, and thos formed with organic acids such as maleic acid, acetic acid, and methanesulfonic

acid.

Bas addition salts includ thos d riv d from inorganic bases, such as ammonium or alkali or alkaline earth metal hydroxides, carbonates, bicarb nat s, and the lik. Such bas s useful in pr paring the salts of this invention thus include sodium hydroxide, potassium hydroxide, ammonium hydroxide, potassium carbonate, s dium carbonate, sodium bicarb nate, potassium bicarbonate, calcium hydroxide, calcium carbonate, and the like. The potassium and sodium salt forms are particularly preferred.

It should be recognized that the particular counterion forming a part of any salt of this invention is not of a critical nature, so long as the salt as a whole is pharmacologically acceptable and as long as the counterion does not contribute undesired qualities to the salt as a whole.

The compounds of the present invention are prepared from compounds of the formula:

The compounds of formula II are defined in Table 1.

TABLE 1
Formula II Compounds^a

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antibiotic	R	R ¹	R ²	R ³	R ⁴	R ⁵	R6	x	Y
vancomycin	н	van	н	инсн3	си ₂ си (си ₃) 2	сн ₂ (со) ин ₂	Н	C1	c:
A82846A	4-epi	4-epi	Н	инсн3	СН ₂ СН (СН ₃) 2	сн ₂ (со) ин ₂	н	н	c 1
A82846B	4-epi	4-epi	н	инсн3	сн ₂ сн (сн ₃) 2	CH2 (CO) NH2	н	C1	C1
A82846C	4-epi	4-epi	н	инсн3	СН ₂ СН (СН ₃) 2	CH2 (CO) NH2	н	н	н
PA-42867-A	4-epi	4-epi	н	инсн3	сн ₂ сн (сн ₃) 2	CH2 (CO) NH2	н	c1	н
PA-42867-C	4-epi	4-epi	н	инсн3	СH2CH(СH3)2	CH2 (CO) NH2	н	H	н
PA-42867-D	4-epi	4-epi	н	N(CH3)2	СН ₂ СН (СН ₃) 2	CH ₂ (CO) NH ₂	н	C1	н
A83850A	н	keto	н	N(CH3)2	СН ₂ СН (СН ₃) 2	CH2 (CO) NH2	н	Cl	C1
A83850B	н	keto	н	NHCH3	СН ₂ СН (СН ₃) 2	CH2 (CO) NH2	н	Cl	c1
actinoidin	actin	acos	н	NH ₂	p-OH,m-Cl-	benzyl	man	CI	н
					phenyl				ļ
avoparcin	risto	risto	man	N(CH3)2	p-rha-	p-0H-	н	н	Н
					phenyl	phenyl			
galacardin	risto	risto	man	мнсн3	p-gal-gal-	p-OH-	н	C1	н
					phenyl	phenyl	[
heleve-	risto	risto	H	инсн3	p-CH ₃ O-rha-	p-OH,m-Cl-	Н	cı	н
cardin			man		phenyl	phenyl			
M47767	actin	acos	Н	инсн3	p-OH,m-Cl-	benzyl	man	Cl	н
					phenyl				

aAbbreviations for the formula II compounds are: actin = actinosaminyl; acos = acosaminyl; 4-epi = 4-epi-vancosaminyl; gal = galactosyl; keto = 4-keto-vancosaminyl; man = mannose; rha = rhamnosyl; rha-gal = rhamnosyl-galactosyl; risto = ristosaminyl; van = vancosaminyl.

In a preferred embodiment of the invention, the formula I compounds are prepared from the A82846 antibiotics (A82846A, A82846B, and A82846C) and PA-42867-A. In a more preferred embodiment, the compounds of the present invention are prepared from A82846B ("A82846B derivatives"). A82846B is represented by formula I compounds wherein R is 4-epi-vancosaminyl, R¹ is hydrogen, R² is NHCH₃, R³ is CH₂CH(CH₃)₂, R⁴ is CH₂(CO)NH₂, R⁵ is hydrogen, R⁶ is 4-epi-vancosaminyl and X and Y are Cl. A82846B derivatives of the present invention having substituents at position R⁵ of formula I are list herein in the manner "R⁵-A82846B". For example, the compound "phenylbenzyl-A82846B" has a phenylbenzyl substituent at position R⁵ in formula

Preferred formula I compounds include those A82846B derivatives wherein R^7 is -(C_1 -Cl₂-alkyl)- R^8 , with -CH₃- R^8 being more preferred, and R^8 is an unsubstituted multicyclic aryl. Of this group, naphthylmethyl-A82846B, acenapthlenyl-methyl-A82846B, and fluorenylmethyl-A82846B are more preferred.

Preferred formula I compounds also include thos A82846B derivativ s wher in R^7 is -(C_1 - C_{12} -alkyl)- R^8 , with - CH_3 - R^8 being more preferred, and R^8 is an unsubstituted h t roaryl r a heteroaryl substituted by halophenyl. Of this group, [1-oxa]fluorenylmethyl-A82846B, chlorophenylbenzoxazolemethyl-A82846B are more preferred.

Further pref rred compounds of formula I include th s A82846B derivativ s wh rein R^7 is -(C_1 - C_{12} -alkyl)- R^8 , with -CH₃- R^8 being mor pref rr d, and R^8 is a group of the formula:

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wherein p is 1 and R^{11} is selected from (C_2-C_5) alkenyloxy, halo- (C_1-C_6) alkoxy, (C_2-C_{10}) alkanoyloxy, (C_1-C_3) alkoxy substituted with (C_1-C_4) alkylthio, and diphenyl- (C_1-C_6) alkyl. Of this group, trifluromethoxybenzyl-A82846B, diphenylmethylbenzyl-A82846B, thiopropylethoxybenzyl-A82846B, acetoxybenzyl-A82846B, non-anoyloxybenzyl-A82846B, and tetrafluoroethoxybenzyl-A82846B are more preferred.

Still further preferred compounds of formula I include those A82846B derivatives wherein R⁷ is -(C₁-C₁₂-alkyl)-R⁸, with -CH₃-R⁸ being more preferred, and R⁸ is a group of the formula:

wherein q is 1 to 5; r is 1; Z is selected from a single bond, divalent (C_1 - C_6)alkyl, divalent (C_2 - C_6)alkenyl, and $-R^{15}$ -($C(R^{14})_2)_8$ -, wherein R^{15} is selected from -O-, -S-, -SO₂-, and -OC(O)-, each R^{14} substituent is hydrogen, and s is 0 or 1; and R^{13} is selected from: (C_4 - C_{10})cycloalkyl; phenyl; and phenyl substituted by nitro, halo, (C_1 - C_{10})alkyl, (C_1 - C_{10})alkoxy, or halo(C_1 - C_3)alkyl. Of this group, chlorophenylbenzyl-A82846B, phenylbenzyl-A82846B, methylphenylbenzyl-A82846B, pentylphenylbenzyl-A82846B, methoxy-phenylbenzyl-A82846B, pentylphenylbenzyl-A82846B, nitrophenylbenzyl-A82846B, phenylbenzyl-A82846B, phenylbenzyl-A82

A82846B, phenyletnynylbenzyl-A82846B, phenoxybenzyl-A82846B, betzylra82846B, butylphenoxybenzyl-A82846B, trifluoromethylphenoxybenzyl-A82846B, dichlorophenoxybenzyl-A82846B, nitrobenzyloxybenzyl-A82846B, benzoyloxybenzyl-A82846B, cyclohexyloxybenzyl-A82846B, cyclohexanoyloxybenzyl-A82846B, thiophenylbenzyl-A82846B, chlorophenylsulfonylbenzyl-A82846B, and cyclohexylbenzyl-A82846B, cyclohexylethoxybenzyl-A82846B, chlorophenoxynitro-benzyl-A82846B, benzylmethoxybenzyl-A82846B, chlorophenoxynitro-benzyl-A82846B, benzoyloxy-dimethoxybenzyl-A82846B, cyclohexanoyloxydimethylbenzyl-A82846B, trifluoromethylphenylbenzyl-A82846B, butylphenylthiobenzyl-A82846B, and bromophenylbenzyl-A82846B more preferred.

Still further preferred compounds of formula I include A82846B derivatives wherein R⁷ is -(C₁-C₁₂-alkyl)-R⁸, with -CH₃- R⁸ being more preferred, and R⁸ is (C₄-C₁₀)cycloalkyl substituted with (C₄-C₁₀)cycloalkyl. Of this group of compounds, more preferred is cyclohexyl-cyclohexylmethyl-A82846B and butylcyclohexylmethyl-A82846B.

Formula I compounds that are prepared from A83850A or A83850B can be prepared from the reduced forms of these compounds. The reduced forms of compounds A83850A or A83850B are produced according to the method described in U.S. Pat. No. 5,187,082, which is incorporated herein by reference.

The compounds of this invention are prepared by reacting a formula il compound with an aldehyde to form an intermediate Schiff's base, which is subsequently reduced with a metal borohydride to give the desired N-alkyl amine.

In the first method of making the compounds of this invention, hereinafter Method A (described in Examples 1 and 2), the reaction for the formation of the Schiff's base is carried out under an inert atmosphere, such as nitrogen or argon, in a polar solvent, such as dimethylformamide (DMF) or methanol (MeOH), or a mixture of polar solvents, such as a mixture of dimethylformamide and methanol, at a temperature of about 25°C to about 100°C. The reaction is preferably carried out at a temperature from about 60°C to about 70°C for 30 minutes to 2 hours in a mixture of dimethylformamide and methanol, or in methanol. The intermediate Schiff's base is then reduced, preferably without isolation, to produce the corresponding N-alkyl derivative(s). The reduction of the Schiff's base can be effected using a chemical reducing agent such as a metal borohydride, for example, sodium borohydride or sodium cyanoborohydrid. The reduction reaction can b carried out in a polar organic solvent, such as dimethylformamid , methanol, or a mixture of polar solvents, such as a mixture of dimethylformamide and m thanol. The r duction reaction can b carried out at a t mperature of about 25°C to about 100°C for 1 to 5 hours. Th r duction reaction is preferably carried out using an excess of sodium cyanobor-

ohydrid in a mixture of dimethylformamide and m thanol or in m thanol at about 60°C to about 70°C for 1 to 2 hours. Method A is preferable for benzylic aldehydes.

In a s cond meth d of making compounds of this invention, her inafter Method B (d scrib d in Exampl 3), the firmation of the Schiff's base is carried out under an inint atmosphere, such as nitrogener argin, in the presence of the reducing agent, sodium cyanoborohydride, in a polar solvent, such as dimethylformamide, methanol, or a mixture of polar solvents, such as a mixture of dimethylformamide and methanol, at a temperature of about 25°C to about 100°C for 1 to 5 hours. The reaction is preferably carried out at a temperature from about 60°C to about 70°C for 1 to 2 hours in a mixture of dimethylformamide and methanol. Method B is preferable for nonbenzylic aldehydes.

In a third method of making compounds of this invention, hereinafter Method C (described in Example 4), the formation of the Schiff's base is carried out a) under an inert atmosphere, such as nitrogen or argon, b) in the presence of the reducing agent, such as a metal borohydride, with sodium cyanoborohydride being most preferred, or a homogenous or heterogeneous catalytic hydrogenation agent(s), such as Crabtree's catalyst, Wilkinson's catalyst, palladium on carbon, platinum on carbon, or rhodium on carbon, c) in a polar solvent, such as dimethylformamide, methanol, or a mixture of polar solvents, such as a mixture of dimethylformamide and methanol, and d) at a temperature of about 25°C to about 100°C. The reaction is preferably carried out at a temperature from about 60°C to about 70°C in methanol. The reaction is allowed to continue for about 20 to about 28 hours, at which time the reaction mixture is adjusted to about pH 7.5 to about pH 10, with a pH of about 9.0 being preferred. The pH adjustment halts the reaction. Because the product is marginally soluble in polar solvents, the solvent of the reaction can be exchanged to an alcohol such as ethanol, butanol, or isopropanol, with isopropanol being preferred, to allow for precipitation of the product. Method C is a preferred method of this invention in view of the increased product yield provided by this method. Another advantage of this reaction scheme is the increased ratio of preferred product (products substituted at the amino group of the sugar denoted as R1 in Formula II compounds) to other products (products that are substituted at the amino groups of substitutents denoted as R and/or R3 of the Formula II compounds). By allowing the reaction to proceed for an extended period of time, such as 20 to 28 hours, products that are monosubstituted at positions denoted as R and R3 in the Formula II compounds are converted to disubstituted forms, making the preferred monosubstituted derivative easier to isolate.

The products of the reaction, obtained from either Method A, B, or C can be purified by preparative reverse-phase HPLC utilizing Waters C18 Nova-Pak columns with ultraviolet light (UV; 235 nm or 280 nm) detection. A 30 minute gradient solvent system consisting of 95% aqueous buffer/5% CH₃CN at time=0 minutes to 20% aqueous buffer/80% CH₃CN at time=30 minutes is typically used, where the aqueous buffer is either TEAP (0.5% aqueous triethylamine adjusted to pH=3 with phosphoric acid) or TFA (0.1% trifluoroacetic acid overall concentration).

HPLC analysis of the reaction mixtures and final purified products can be accomplished utilizing a Waters C18 MicroBonda-Pak column (typically 3.9 x 300 mm steel) or Waters Nova-pak C18 RCM column (8 x 100 mm) with UV (235 nm or 280 nm) detection. A 30 minute gradient solvent system consisting of 95% aqueous buffer/5% CH₃CN at time=0 minute to 20% aqueous buffer/80% CH₃CN at time=30 minutes is typically used, where the aqueous buffer is either TEAP (0.5% aqueous triethylamine adjusted to pH=3 with phosphoric acid) or TFA (0.1% trifluoroacetic acid overall concentration).

The ratio of the aldehyde to the formula II compound and the reaction conditions determines the products of the reaction. The monosubstituted derivatives are those derivatives where a hydrogen atom of the amino group at position R¹ in formula II is replaced by one of the substituents listed above for formula I. When using Methods A or B, described above, the formation of monosubstituted derivatives substituted at the amino group of the amino sugar at position R¹ in the formula II compounds is favored by using a slight excess of aldehyde, a shorter reaction time, and a lower temperature. As noted above, Method C favors the formation of the monosubstituted derivative. The monosubstituted derivative is preferred. A large excess of the aldehyde favors the formation of disubstituted and trisubstituted derivatives of the formula II compounds. The disubstituted derivatives are the derivatives where a hydrogen atom at two of the locations selected from the amino group at position R³ and the amino group of the amino sugars designated as R or R¹ in formula II, are replaced by the reduced aldehyde moiety. The trisubstituted derivatives are the derivatives where a hydrogen atom at three of the locations selected from the amino group at position R³ and the amino group of the amino sugars designated as R or R¹ in formula II, are replaced by the reduced aldehyde moiety.

Examples of compounds that have been prepared and are illustrative of the formula I compounds are listed in Tabl s 2A and 2B. Table 2A lists compounds pr pared by reacting an ald hyde with the glycop ptide A82846B. Table 2A lists the sidechain substitutions on the amino group of the 4- pi-vancosaminyl sugar of the 4-epi-vancosaminyl-O-glycosyl disaccharide of the A82846B compound. All of the compounds listed are monosubstituted derivatives.

Tabl 2B lists thos compounds that were prepared by reacting an aldehyde with a varity of glycopeptid antibitics oth rithan A82846B. The compounds of Tabl 2B are monosubstituted at the amino group of the amino sugar disignated as R1 in firmula II with the sid chain list d. All fithe compounds listed are minosubstituted derivatives.

TABLE 2A

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	COMPOUND NO.	SIDECHAIN
	1	2-naphthylmethyl
10	2	4-phenylbenzyl
	3	1-naphthylmethyl
	4	4-phenoxybenzyl
45	5	4-benzyloxybenzyl
15	6	4-trifluoromethoxybenzyl
	7	4-allyloxylbenzyl
	8	4-nonyloxybenzyl
20	9	2-methoxy-1-naphthylmethyl
	10	4-dodecyloxybenzyl
	11	9-phenanthranylmethyl
	12	4-decyloxybenzyl
25	13	9-anthranylmethyl
	14	4-[phenylethynyl]4-phenylbenzyl
	15	4-methoxy-1-naphthylmethyl
30	16	1-pyrenylmethyl
•	17	9-[10-methyl]anthranylmethyl
	18	9-[10-chloro]anthranylmethyl
	19	2-benzthienylmethyl
35	20	4-[4-hydroxyphenyl]benzyl
	21	4-[4-octylphenyl]benzyl
	22	4-[4-pentylphenyl]benzyl
	23	4-[4-octyloxyphenyl]benzyl
40	24	3-pyridylmethyl
	25	5-nitro-1-naphthylmethyl
	26	4-pyridylmethyl
45	27	4-quinolylmethyl
	28	3-quinolylmethyl
	29	4-stilbenzyl
	30	2-quinolylmethyl
50	31	2-pyridylmethyl
	32	2-fluorenylmethyl
	33	4-phenoxyphenethyl

TABLE 2A

COMPOUND NO.	SIDECHAIN
34	4-[4-pentylcyclohexyl]benzyl
35	4-benzylphenethyl
36	4-[4-biphenyl]benzyl
37	4-trifluoromethylbenzyl
38	trans-cinnamyl
39	4-[1-oxa]fluorenylmethyl
40	4-[4-pentoxyphenyl]benzyl
41	4-thiomethylbenzyl
42	2,3-[2-methyl-3-[4-t-butylphenyl]]propenyl
43	9-(1-methyl)-acridinylmethyl
44	2-hydroxy-1-naphthylmethyl
45	4-{2-phenyl-6-methoxy quinoy methyl
46	4-diphenylmethylbenzyl
47	3,4 cyclohexenylmethyl
48	3,4-methylenedioxylbenzyl
49	3-phenoxybenzyl
50	4-benzylbenzyl
51	3-benzyloxy-6-methoxy benzyl
52	4-benzyloxy-3-methoxybenzyl
53	3,4-dibenzyloxybenzyl
54	4-[4-methoxyphenyl]benzyl
55	4-[3-cyanopropoxy]benzyl
56	3,4-ethylenedioxybenzyl
57	4-[4-nitrophenoxy]benzyl
58	2,3-methylenedioxybenzyl
59	2-benzyloxyphenethyl
60	2-ethoxy-1-naphthylmethyl
61	2-benzylfurylmethyl
62	3-phenoxyphenethy1
63	4-phenoxyphenethy1
64	4-[4-nitrophenyl]benzyl
65	5-methoxy-2-naphthylmethyl

TABLE 2A

COMPOUND NO.	SIDECHAIN
67	5-phenyl-2-thienylmethyl
68	4-benzyloxyphenethyl
69	3-benzyloxyphenethyl
70	4-[2-nitrophenoxy]benzyl
71	5-[4-methoxyphenyl]-2-thienylmethyl
72	4-difluormethoxybenzyl
73	2,3,4,5,6-pentamethylbenzyl
74	5-iodo-2-thienylmethyl
75	4-[2-[2-chloroethoxy]ethoxy]benzyl
76	3,4-dimethylbenzyl
77	3-acetoxybenzyl
78	4-nitrobenzyl
79	4-phenylethynylbenzyl
80	4-[2-chloro-6-fluorobenzyloxy]benzyl
81	4-[3,4-dichlorophenoxy]benzyl
82	5-[2,3-dihydrobenzfuryl]methyl
83	4-[2-(N, N-diethylamino) ethoxy]benzyl
84	2-bicyclo[2.1.2]heptylmethyl
85	2-hydroxy-5-phenylbenzyl
86	3-[4-chlorophenoxy]benzyl
87	4-[3-chlorophenoxy]-3-nitrobenzyl
88	4-[2-chlorophenoxy]-3-nitrobenzyl
89	3,5-dimethylbenzyl
90	4-[4-ethylphenyl]benzyl
91	3-phenylbenzyl
92	4-[3-fluorophenyl]benzyl
93	4-[4-chlorobenzyloxy]benzyl
94	4-[4-chlorophenoxy]-3-nitrobenzyl
95	4-[4-methylphenoxy]benzyl
96	4-{4-t-butylphenoxy}benzyl
97	4-[4-methylphenyl]benzyl
98	4-[4-methoxyphenoxy]benzyl
59	4-acetoxy-3-methoxybenzyl

TABLE 2A

COMPOUND NO.	SIDECHAIN
100	4-[(2-phenyl)ethyl]benzyl
101	3-[5-phenyl]pyridinylmethyl
102	4-[2-nitrophenyl]benzyl
103	2-[1-hydroxy]fluorenylmethyl
104	4-benzyl-3-methoxybenzyl
105	4-[cyclohexylmethoxy]-3-ethoxybenzyl
106	3-[3,3'-dichlorophenoxy]benzyl
107	4-[4-propylphenyl]benzyl
108	4-thiophenylbenzyl
109	4-[alpha-hydroxybenzyl]benzyl
110	2,2-dinitro-4-thiophenebenzyl
111	3-[3-trifluoromethylphenoxy]benzyl
112	4-[t-butylethynyl]benzyl
113	4-phenoxy-3-methoxy-benzy1
114	4-[3-trifluoromethylphenoxy]-3-nitrobenzyl
115	2-phenylthiobenzyl
116	2-[4-chlorophenyl]-6-benzoxazolemethyl
117	4-[alpha-methoxybenzyl]benzyl
118	4-cyclohexylbenzyl
119	3-[3,4-dichlorophenoxy]benzyl
120	acenaphthlenylmethyl
121	4-{1,1,2,2-tetrafluoroethoxy}benzyl
122	4-benzoyloxy-3,3'-dimethoxybenzyl
123	3-[cyclohexylmethoxy]benzyl
124	4-cyclohexyloxybenzyl
125	3-[2-quinoylmethoxy]benzyl
126	4-[alpha-ethoxybenzyl]benzyl
127	4-{cyclohexylethoxy benzyl
128	4-{alpha-propoxybenzyl]benzyl
129	4-[4-methyl-1-piperidino]benzyl
130	2-thiophene-1,2-cyclohexenylmethyl
131	4-[4-nitrobenzyloxy]benzyl
132	3-[4-trifluoromethylphenoxy]benzyl

TABLE 2A

COMPOUND NO.	SIDECHAIN
133	3-benzoyl-2,4-dichlorobenzyl
134	4~[2-(2-thiopropyl)ethoxy]benzyl
135	4-[2-methyl-1-piperidino]benzyl
136	4-hydroxybenzyl
137	4-{2-pyridyl}benzyl
138	4-acetoxybenzyl
139	5,6-benzonorbornylmethyl
140	3-phenylcyclopentylmethyl
141	1-adamantylmethyl
142	3-[cyclohexylmethoxy]-4-methoxybenzyl
143	2-[2-glucosyl]benzyl
144	4-[4-pentoxybiphenyl]benzyl
145	3,4-dihydroxybenzyl
146	4-[4-methylpiperazino]benzyl
147	4-morpholinobenzyl
148	4-[4-chlorophenylsulfonyl]benzyl
149	4-methylsulfonyloxybenzyl
150	4-benzoyloxybenzyl
151	5-phenyl-3-pyridinylmethyl
152	<pre>4-{N,N-bis(2-chloroethyl)amino}benzyl</pre>
153	3-cyclohexyloxybenzyl
154	4-[2-t-butoxyethoxy]benzyl
155	3,3'-dichloro-4-hydroxy-benzyl
156	1,2,3,4,-tetrahydro-9-anthranylmethyl
157	4-cyclohexanoyloxybenzyl
158	4-nonanoyloxybenzyl
159	4-[phenylsulfinyl]benzyl
160	4-anilinobenzy1
161	cyclohexylmethyl
162	3-benzoyloxybenzyl
163	3-nonanoyloxybenzyl
164	4-[cyclohexyl]cyclohexylmethyl
165	3-cyclohexanoyloxybenzyl

TABLE 2A

C	OMPOUND NO.	SIDECHAIN
	166	4-[cyclohexanoyloxy]-3,3'-[dimethoxy]benzyl
	167	4-[nonanoyloxy]-3,3'-[dimethoxy]benzyl
	168	1,2,3,4-tetrahydro-6-naphthylmethyl
	169	2-hydroxybenzyl
	170	[2-[6,6-dimethyl-bicyclo[3.1.1]hept-2-enyl]methyl
	171	1-cyclohexenyl-4-isopropylmethyl
-	172	4-[4-methoxyphenyl]butyl
<u> </u>	173	4-[[2,3,4,5,6-pentamethyl]phenylsulfonyloxy]benzyl
	174	4-[1-pyrrolidinosulfonyl]benzyl
	175	3-[4-methoxyphenyl]propyl
	176	8-phenyloctyl
	177	4-[2,3-dihydroxypropoxy]benzyl
	178	4-{N-methylanilino}benzyl
	179	2-[2-napthyl]ethyl
	189	6-methyl-2-naphthylmethyl
-	190	cis-bicyclo[3.3.0]octane-2-methyl
	191	2-tridecynyl
	192	4-butyl-2-cyclohexylmethyl
	193	4-[(4-fluorobenzoyl)amino]benzyl
	194	4-[(3-fluorobenzoyl)amino]benzyl
	195	8-phenoxyoctyl
	196	6-phenylhexyl
	197	10-phenyldecyl
	198	8-bromooctyl
	199	11-tridecynyl
	200	8-[4-methoxyphenoxy]octyl
	201	8-[4-phenylphenoxy]octyl
	202	8-[4-phenoxyphenoxy]octyl
-	203	3-[3-trifluoromethylphenoxy]benzyl
	204	10-undecenyl
	205	4-cyclohexylbutyl
	206	4-phenyl-2-fluorobenzyl
	207	^-hexadecynyl

TABLE 2A

COMPOUND NO.	SIDECHAIN
208	3-{cyclopentyl}propyl
209	4-[2-methylphenyl]benzyl
210	4-[phenylazo]benzyl
211	4-{4-flurophenyl]benzyl
212	3-nitro-4-[4-nitrophenyl]benzyl
213	3-nitro-4-[2-nitrophenyl]benzyl
214	9-decenyl
215	4-[3,4-dimethoxyphenyl]benzyl
216	4-[4-trifluromethylphenyl]benzyl
217	5-hexenyl
218	4-{2-thienyl}benzyl
219	4-[6-phenylhexyloxy]benzyl
220	9,10-dihydro-2-phenantrene methyl
221	4-[3,4-dimethylphenyl]benzyl
222	4-[4-methylphenyl]-2-methylbenzyl
223	4-[3-phenylpropyloxy]benzyl
224	4-[3-methylphenyl]benzyl
225	4-(4-methylphenyl)-3-methylbenzyl
226	4-[4-pentenyloxy]benzyl
227	4-[1-heptynyl]benzyl
228	3-[4-t-butyl-phenylthio]benzyl
229	4-[4-chlorophenyl]benzyl
230	4-[4-bromophenyl]benzyl
231	4-[4-cyanophenyl]benzyl
232	4-[1-nonynyl]benzyl
233	4-[11-tridecynyloxy]benzyl
234	12-phenyldodecyl
235	6-phenyl-5-hexynyl
236	11-phenyl-10-undecynyl
237	4-[2-methylphenyl]-3-methylbenzyl
238	3-[2'-thienyl]-2-thienylmethyl
239	4-[benzyloxymethyl]cyclohexylmethyl
240	4-[4-chlorophenoxy]benzyl

TABLE 2A

5	COMPOUND NO.	SIDECHAIN	
	241	4-{benzyl}cyclohexylmethyl	
	242	4-benzoylbenzyl	
10	243	4-{phenoxymethyl}benzyl	
	244	4-[4-chlorobenzyl]benzyl	

TABLE 2B

COMPOUND NO.	GLYCOPEPTIDE CORE	SIDECHAIN
180	vancomycin	1-napthylmethyl
181	vancomycin	4-phenylbenzyl
182	A82846A	4-phenylbenzyl
183	A82846C	4-phenylbenzyl
184	A82846C	4-phenoxybenzyl
185	PA-42867 A	4-phenylbenzyl
186	reduced A838450A	4-phenylbenzyl
187	alpha-avoparcin	4-phenylbenzyl
188	beta-avoparcin	4-phenylbenzyl

The formula I compounds have <u>in vitro</u> and <u>in vivo</u> activity against Gram-positive pathogenic bacteria. The minimal inhibitory concentrations (MIC) at which the formula I compounds inhibit certain bacteria are given in Table 3. The MIC's were determined using a standard broth micro-dilution assay.

Organism	vancomycin A82846A		A82846B	A82846C	1	2	3	4	5	9
Staphylococcus aureus 446	0.5	0.25	0.25	0.5	≥0.06	≥0.05	≥0.05	≥0.06	1	0.5
Staphylococcus aureus 489	0.125	0.5	≥.06	≥.06	• •	0.25	≥0.06	\$0.06	0.5	0.25
Staphylococcus aureus 447	0.5	0.25	0.25	0.5	≥0.06	≥0.05	S0.06	0.25	0.5	5.0
Staphylococcus aureus X400	0.5	0.125	0.125	0.25	20.06	1	0	20.06		-
Staphylococcus aureus X778	0.5	0.125	0.125	0.5	~	≥0.06	≥0.06	20.06	0.5	0.25
Staphylococcus aureus 491	1	0.25	0.25	1	NI	≥0.06	0.5	\$0.06	0.5	0.125
Staphylococcus aureus S13E	0.5	0.125	0.125	0.25	0.125	≥0.06	50.05	≥0.06	1	0.25
Staphylococcus aureus SA1199	0.5	0.125	0.125	0.25	≥0.06	0.5	0.125	≥0.06	1	0.25
Staphylococcus aureus SA1199A	0.125	≥.06	≥.06	0.125	≥0.06	≥0.06	50.06	≥0.06	\$0.06	≥0.06
_	0.5	S.06	0.125	S.06	'	20.06	0	20.06	\$0.06	
Staphylococcus haemolyticus 105	16	0.5	1	-4	7	~	4	0.5	7	5.0
Staphylococcus haemolyticus 415	8	1	4	2	4		60	0.5		0.5
Staphy lococcus epidermidis 270	16	0.25	0.25	0.125	80	œ	œ	\$0.06	0.25	0.125
Entercoccus faecium 180	>64	16	æ	16	0.5	0.25	0.5	0.125	20.06	0.125
Entercoccus faecium 180-1	0.5	0.125	0.125	0.125	so.06	50.06	\$0.06	\$0.06	20.06	\$0.08
Entercoccus faecalis 2041	7	0.125	0.25	0.5	0.125	0.125	80.06	\$0.06	\$0.06	20.06
Entercoccus faecalis 276		0.125	0.125	0.5	≥0.06	0.5		\$0.06	0,	\$0.06
Entercoccus gallinarum 245	5	0.125	0.25	0.5		\$0.06	20.06	50.06		≥0.06
Haemophilus influenzae RD	>64	>64	>64	>64	>64		, , ,			>64
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	× 64	>64
Streptococcus pyogenes C203	0.5			0.125	50.06	\$0.06	\$0.06	\$0.06	\$0.06	\$0.06
Streptococcus pneumoniae Pl	0.25			>.06	≥0.06	50.06	\$0.06	50.08	\$0.06	\$0.06

Organism	7	8	6	10	11	12	13	14	15	16	17
Staphylococcus aureus 446	8	2	2	16	4	32	2	4	1	4	71
Staphylococcus aureus 489	7	4	0.5	>64		80	1	7	≥0.06	0.5	٦.
Staphylococcus aureus 447	3	80	4	>64	4	32	æ	80	7	4	æ
Staphylococcus aureus X400	1	8	0.5	>64	0.5	80	-	4	0.25	0.5	0.5
Staphylococcus aureus X778	0.25	œ	0.25	16	0.25	80	2	4	0.25	7	0.5
Staphylococcus aureus 491	2	4	0.5	91	7	4	7		0.25	٦	~
Staphylococcus aureus S13E	7	80	0.5	8	0.5	æ	0.25	4	0.5		
Staphylococcus aureus SA1199	7	2	0.25	80	2	00	0.5	80	0.25	7	₹ :
9	\$0.06	2	\$0.06	4	≥0.06	0 0	≥0.05	0.5	≥0.06	20.06	\$0.06
	1		0.25	8	7		4	8	0.25	-	7
	8	æ	4	>64	4	16	8	4	0.5	80	æ
Staphylococcus haemolyticus 415	91	80	4	>64	7	32	1	æ	7	4	80
Staphylococcus epidermidis 270	4	4	16	>64	2	0.125	æ	4	-	7	9
Entercoccus faecium 180	7	1	1	8	1	4	2	1	0.5		~
Entercoccus faecium 180-1	S 0.06	0.5	≥0.06	4	≥0.06	7	20.06	1	\$0.06	0.125	20.06
Entercoccus faecalis 2041	_	9	0.25	16	0.5	91	0.125	~1 ·	ě.	0	0.25
Entercoccus faecalis 276	-	¥	0.26	18	1	7	0.5	4	20.06	~	5.0
Entercoccus gallinarum 245	5.0	ao	0.25	œ	≥0.06	32	0.25	0.25	\$0.06		0.5
Haemophilus influenzae RD	16	>64	≥0.06			64	32		1	4	32
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	9 9	>64	>64
Streptococcus pyogenes C203	50.05	≥0.06	\$0.08	0.5	50.06	0.25	S0.06	50.06	≥0.06	20.06	20.06
Streptococcus pneumoniae P1	\$0.08	S Ø.06	80.08	0.125	50.06	20.06	\$0.06	50.06	50.06	50.06	50.06

Organism	18	19	20	21	22	23	24	25	26	27	28
Staphylococcus aureus 446	2	0.5	0.5	>64	16	38	0.5	0.5	0.25	7	0.25
Staphylococcus aureus 489	п	0.25	0.5	32	8	>64	≥0.06	\$0.06	S 0.06	20.06	\$0.0 6
Staphylococcus aureus 447	80	-	4	>64	16	16	-1	0.25	2	80	-
Staphylococcus aureus X400	П	0.25	9.0	32	æ	16	0.25	≥0.06	0.25	0.5	\$0.06
Staphylococcus aureus X778	0.5	0.25	0.25	32	æ	16	0.125	S0.06	0.125	0.5	20.06
Staphylococcus aureus 491	2	2	1	64	æ	16	0.5	0.125	0.5	-	0.25
Staphylococcus aureus S13E	1	≥0.06	20.06	64	16	16	≥0.06	≥0.06	0.25	0.125	₹0.06
Staphylococcus aureus SA1199	7	0.5	2	64	16	16	0.5	≥0.06	-	0.5	0.125
Staphylococcus aureus SA1199A	\$0.06	20.06	S0.05	16	7	16	≥0.05	≥0.06	≥0.06	20.06	20.06
Staphylococcus aureus SA1199B	7		0.5	64	16	16	7	0.125	0.5		0.125
Staphylococcus haemolyticus 105	16	4	8	>64	16	4	4	-1	4	16	♥:
Staphylococcus haemolyticus 415	8	80	4	64	16	16	≥0.06	32	8	œ	au :
Staphylococcus epidermidis 270	æ	2	2	32	4	64	7	0.5		4	، اسم
Entercoccus faecium 180	2	Н	1	8	-	>64	4	0.5	4	30	
Entercoccus faecium 180-1	20.06	S0.06	90.0≥	8	20.06	32	S0.06	50.06	0.25	0.5	\$0.06
Entercoccus faecalis 2041	0.25	≥0.06	≥0.05	32	2	32	≥0.06	0.25	0.25	0.125	0.25
Entercoccus faecalis 276	1	≥0.06	0.25	64	49	32	0.25	0.25	\$0.06	0.5	20.06
Entercoccus gallinarum 245		20.06	0.25	æ	-	œ:	0.25	≥0.06	0.125	0.5	0.25
Haemophilus influenzae RD	16	32	æ	>64	64	>64	>64	32	>64	>64	>64
Escherichia coli EC14	>64	>64	>64	>64	>64	×64	>64	>64	>64	^64	>64
Streptococcus pyogenes C203	50.06	≥0.06	≤0.06	7	S0.06		50.06	20.06	50.06	20.06	
Streptococcus pneumoniae Pl	S0.06	\$0.06	\$0.06	0.5	0.25	0.5	\$0.06	\$0.06	S0.06	20.06	

Organism	29	30	31	32	33	34	35	36	37	38	39
Staphylococcus aureus 446	ī	1	0.5	1	4	32	0.5	8	0.5	9.0	0.125
Staphylococcus aureus 489	1	0.125	≥0.06	1	≥0.06	æ	≥0.06	2		≥0.06	≥0.06
Staphylococcus aureus 447	0.25	7	0.5	0.5	0.125	œ	0.125	7	0.125	0.125	0.25
Staphylococcus aureus X400	0.25	≥0.06	0.125	0.5	0.25	32	0.25	4	0.25		30.05
Staphylococcus aureus X778	90.05	≥0.06	0.125	0.5	0.5	16	≥0.0€	7	≥0.0€	0.5	\$0.05
Staphylococcus aureus 491	0.25	0.5	0.5	0.25	0.125	8	0.125	1	0.25	0.5	0.25
Staphylococcus aureus S13E	-	0.125	0.25	1	0	16	90.0≥	7	≥0.06	≥0.06	\$0.05
Staphylococcus aureus SA1199	0.25	0.5	0.25	1	П	16	0.25	4	0.25	-	≥0.06
Staph/lococcus aureus SA1199A	50.05	≥0.06	≥0.06	90.0≥	≥0.05	7	20.06	≥0.06	≥0.06	0.0	\$0.06
Staphylococcus aureus SA1199B	0.25	0.125	0.25	0.125		16	0.25	4	20.06	0.125	20.06
Staphylococcus haemolyticus 105	4	4	7	4		32	7	4	0.25		7
S	~	16	16	7	20	>64	4	∞	7	-	4
Staphylococcus epidermidis 270	0.5		1	1	7	16			0.25	0.5	0.25
	0.25	2	4	0.25	7	4	-	0.25	0.125	50.06	0.5
İ	20.06	S	\$0.05	≥0.06	\$0.06	~	≥0.06	S0.06	•	\$0.06	≥0.06
	0.25		0.25	0.25	S0.06	œ	S0.06			20.06	\$0.06
Entercoccus faecalis 276	0.25		0.25	0.125	20.06	16	≥0.06	7		0.5	\$0.06
45	0.25		0.25	0.25	0.25	4	\$0.06	0.25	0.125	0.125	\$0.06
Haemophilus influenzae RD	>64		>64	>64						:	
Escherichia coli EC14	99	>64	>64	32	>64	>64	>64	>64	>64	>64	· • • •
Streptococcus pyogenes C203									50.06	≥0.06	50.06
Streptococcus pneumoniae P1		,							\$0.08	50.06	S0.06

Organism	40	41	42	43	44	45	46	47	48	49	50
S 44	4	2	1	0.5	0.25	1	1	0.125	0.125	0.5	0.5
Staphylococcus aureus 489	7	20.06	0.5	≥0.06	≥0.06	0.5	1	≥0.06	≥0.06	≥0.06	≥0.06
Staphylococcus aureus 447	2	0.25	0.5	2	-	16	7	2	2	ا ـــا	0.5
Staphylococcus aureus X400	7	\$0.06	1	0.25	≥0.06	0.25	2	≥0.06	≥0.06	0.125	0.125
Staphylococcus aureus X778	4	0.125	1	50.06	S0.06	0.25	2	50.05	20.06	0	
Staphylococcus aureus 491	4	0.5	0.5	1	0.125	-	2	0.5	0.25	0.125	0.5
Staphylococcus aureus S13E	4	≥0.06	0.5	0.25	0.25	0.5	7	≥0.06	≥0.06	0.	0.125
Staphylococcus aureus SA1199	4	\$0.06	1	0.5	0.25	2	2	0.5	0.25	~	
Staply lococcus aureus SA1199A	0.5	S0.06	90.0≥	≥0.06	≥0.06	≥0.06	0.5	0.25	\$0.06	50.06	≥0.0€
Staphylococcus aureus SA1199B	80	0.25	2	0.5	0.25		7	0.25	7		2
Staphylococcus haemolyticus 105	2	2	2	Þ	2	16	2	4	2	1	0.5
Staphylococcus haemolyticus 415	2	4	1	8	4	8	2	16	æ	-	7
Staphylococcus epidermidis 270	1	0.25	9.0	2	0.5	80	2	1	1	7	0.5
Entercoccus faecium 180	1	0.25	0.25	4	80	-	0.5	2	_	0.25	0.25
Entercoccus faecium 180-1	2	≥0.06	≥0.06	≥0.06	≥0.06	\sim 1	\$0.06	≥0.06	≥0.06	≥0.06	\$0.06
Entercoccus faecalis 2041	-	≥0.06	0.125	0.5	\$0.06	0.125	-	50.06	\$0.06	0;	≥0.06
Entercoccus faecalis 276	7	20.06	8	0.5	0.125	0.25	0.5	50.06	\$0.06	7	0.25
Entercoccus gallinarum 245	11	≥0.06	-	0.5	0.5	0.5	0.25	16	1		7
Haemophilus influenzae RD					>64	>64	>64	>64	>64	>64	>64
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	\$0.06	50.06	≥0.06	≥0.06			١ ١				
Streptococcus pheumoniae Pl	<0.05	90°05	90.08	<0.05		90.0 ×	>0.06	≥0.06	≥0.06	S0.06	≥0.06

Organism	51	52	53	54	55	99	57	58	59	9	61
Staphylococcus aureus 446	0.25	≥0.06	2	1	0.5	0.5	0.25	0.25	0.5	1	0.5
Staphylococcus aureus 489	≥0.06	0.5	2	20.06	-		0.5	≥0.06	0.125	0.5	7
Staphylococcus aureus 447	0.5	≥0.06	4	0.25	4	7	0.5	1	-	2	7
Staphylococcus aureus X400	\$0.05	≥0.06	4	\$0.06	S0.06	≥0.06	0.125	\$0.05	0.25	0.5	≥0.06
Staphylococcus aureus X778	0.5	0.5	7	\$0.06	0.5	0.125	≥0.06	\$0.06	≥0.06	0.25	0.125
Staphylococcus aureus 491	0.25		~		0.5	0.5		0.125			0.5
Staphylococcus aureus S13E	٠.	0.5	2	0.5	0.5	0.125	S0.06	≥0.06	0.125	0.25	0.125
Staphylococcus aureus SA1199	0.5	2	2	0.5	0.5	0.5	П	-	S0.06	0.5	0.25
Staphylococcus aureus SA1199A	\$0.06	50.08	-	S0.06	20.06	S0.06	≤0.06	S0.06	\$0.06	50.06	50.06
Staphylococcus aureus SA1199B	_	2	2	7	0.5	0.5	0.125	0.125	0.5	0.5	0.25
Staphylococcus haemolyticus 105	0.5	0.5	2	2	4	4	8	7	80	>64	64
Staphylococcus haemolyticus 415	1	1	2	1	16	91	-	80	œ	16	80 :
Staphylococcus epidermidis 270	0.5	0.5	2	0.25	-		0.5	4	-	C	-
Entercoccus faecium 180	0.5	7	1	1	2	7	0.5	œ	80	7	7
Entercoccus faecium 180-1	20.06	≥0.06	2	20.05	≥0.06	≥0.06	≥0.06	0.25	≥0.06	≥0.06	≥0.06
Entercoccus faecalis 2041	20.06	0.5	1	≥0.0€	0.125	0.25	≥0.06	0.5	0.5	0.25	20.06
Entercoccus faecalis 276	20.06	0.125	80	1	0.5	0.25	0.5	0.5	0.125	0.5	0.25
Entercoccus gallinarum 245	-	7	7	0.5	16	16	7	0.5		16	œ :
Haemophilus influenzae RD	>64	>64	>64		>64	>64	>64	>64	>64	>64	>64
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203		≥0.06	≥0.06	50.06	\$0.06	≥0.06	S0.06	\$0.06	20.06		
Streptococcus pneumoniae P1	≥0.06	S0.08	\$0.06	\$0.06	≥0.06	≥0.06	30.05	≥0.06	20.06	≥0.06	≥0.06

TABLE 3
In Vitro Activity of Formula I Compounds
MIC (mcg/ml)/Compound

Organism	62	63	64	65	99	67	68	69	7.0	7.1	72
Staphylococcus aureus 446	2	0.5	0.25	2	0.25	0.25	0.125	1	0.125	4	2
Staphylococcus aureus 489	7	æ	0.25	0.125		≥0.06	0.125	0.25	\$0.0¢	0.25	\$0.06
Staphylococcus aureus 447	0.5		0.5		-1		0.25		0.5	4	-
Staphylococcus aureus X400	20.06	20.06	0.125	0.125	0.125		0	S	≥0.06	-	0.125
Staphylococcur aureus X778	0.5	0.125	7	0.5	≥0.06	0.25	0	-	20.06	7	0.25
Staphylococ aureus 491	0.125	0.5	0.125	0.5	0.25		2		0.5	2	0.25
Staphylococcus aureus S13E	0.5	0.125	2	0.5	≥0.06	0.25	≥0.06	0.25	≥0.06		≥0.06
Staphylococcus aureus SA1199	0	0.25		0.5	0.25		0	-	≥0.06	· =4	
Staphy lococcus aureus SA1199A	\$0.06	0.125	≥0.06	\$0.06	\$0.08	≥0.06	0	≥0.06	≥0.06	0.25	۰.
Staphylococcus aureus SA1199B		0.5	0.125	2	0.25	-	S	~	≥0.06		≥0.06
Staphylococcus haemolyticus 105	7	7	64	64	64	64	2	7	7	16	
Staphylococcus haemolyticus 415	4	æ	2	7	œ (2	4	æ	7	00	• ❤
Staphylococcus epidermidis 270	-	1	0.5			0.5	2	7	0.25		0.25
Entercoccus faecium 180	4	16	0.125	0.5	7	0.25	2	4			9.0
Entercoccus faecium 180-1	\$0.06	≤0.06	≥0.06	≥0.06	90.	≥0.06	≥0.06	50.06	≥0.06	0.25	≥0.06
Entercoccus faecalis 2041	-	0.25	≥0.06	≥0.05	9	S0.06	20.06	0.25	0	: 🗕	≥0.06
Entercoccus faecalis 276	0.5	0.5	•	0.5	90.	≥0.06	0	0.5	0.0	. ~	\$0.08
Entercoccus gallinarum 245	49 1	001	2	7	60	2	4	, c c	~	: : œ	4
Haemophilus influenzae RD	>64	>64	>64	>64	^	9	· vo	>64	16	>64	32
Escherichia coli EC14	>64	>64	>64	>64	>64	· 64	>64	>64	×64	799	>64
Streptococcus pyogenes C203	\$0.06	20.06		: !			ŀ		:	:	
Streptococcus pneumoniae P1	<0.0>	90.0	90 OS	40 0×	20 08	×0 0×				! !	

TABLE 3
In Vitro Activity of Formula I Compounds
MIC (mcg/ml)/Compound

Organism	73	74	75	76	77	7.8	79	80	81	82	83
Staphylococcus aureus 446	0.25	Þ	2	0.25	30.05	2	2	4	7	2	:
Staphylococcus aureus 489	0.25	≥0.06	20.06	≥0.06	≥0.06	≥0.06	2	7	7	0.25	0.25
Staphylococcus aureus 447			-1	0.5	7	20.06	2	2	~	7	C3:
Staphylococcus aureus X400	0.5	≥0.06	\$0.06	0.25	≥0.06	≥0.06	0.25	4	-	0.25	2
Staphylococcus aureus X778	1	20.06	50.05	0.25	≥0.06	\$0.06	7	0.5	,q.	0.5	0.5
Staphylococcus aureus 491	0.25	0.125	0.25	0.25		0.25	4	1	1		0.5
Staphylococcus aureus S13E		0	0		0.125		4		0.5	≥0.06	0.125
Staphylococcus aureus SA1199	0.5	\$0.06	2	S 0.06	≥0.06	0.125	-	2	7	0.25	7
Staphylococcus aureus SA1199A	0.25	≥0.06	≥0.06	0.125	S0.06	≥0.06	0.125	1	0.5	0.5	0.25
Staphylococcus aureus SA1199B	\$0.06	-	0.5	0.25	0.125	0	-	1	-	-	
Staphylococcus haemolyticus 105	0.5	4	2	2	2	7	4	7	-4	80	ر ان
Staphylococcus haemolyticus 415	2	4	4	4	80	16	4	4	7	80	7
Staphylococcus epidermidis 270	0.125	0.5	0.5	0.25	0.5	0.5	0.5	2	1	4	7
Entercoccus faecium 180	0.5	0.5	0.5	0.5	89	-	≥0.06	0.125	0.0	2	æ;
Entercoccus faecium 180-1	≥0.06	≥0.05	\$0.06	≥0.06	0.125	≥0.06	50.06	≥0.06	0.125	0.125	0.125
Entercoccus faecalis 2041	0.125	≥0.06	\$0.06	\$0.06	0.25	≥0.0€	20.06	20.06	0:	φ:	0.5
Entercoccus faecalis 276	0.25		≥0.06	\$0.05	0.25	0.125	S0.06	50.06	\$0.06	0	:
Entercoccus gallinarum 245	7	\$0.06	4	4	0.25	12	20.06	\$0.06	0.25	0.125	0.5
Haemophilus influenzae RD	0.25	0.5	2	>64	64		16	16	16	64	>64
Escherichia coli EC14	>64	>64	>64	>64	>64	×64	>64	>64	>64	>64	764
Streptococcus pyogenes C203	\$0.06	≥0.06	\$0.08	20.06	50.06	\$0.06	20.06	50.06	S0.06	\$0.06	\$0.06
Streptococcus pneumoniae P1	\$0.06	20.06	50.08	20.06	20.06	50.06	20.06	20.06	\$0.06	50.06	\$0.06

Ordanism	84	85	98	87	88	89	06	91	92	93	9.4
Staphylococcus aureus 446	0.5	0.125	1	1		0.5	2	S	1''	2	1
Staphy lococcus aureus 489	S0.06	0.25	1	0.5	0.5	0.25	7	≥0.06	≥0.06	0.25	. ~
Staphylococcus aureus 447	4	0.125	0.5	0.5			-	S	S	0.25	0.5
Staphylococcus aureus X400	≥0.06	0.25	1	1	≥0.06	0.25	-	S	0.5	≥0.06	-
Staphylococcus aureus X778	≥0.06	0.25		2	0.5	0.25	-	≥0.06	•	-	0.5
Staphylococcus aureus 491	-	0.125	7	2	0.5	. ⊶	2	-	7	0.25	0.5
Staphylococcus aureus S13E	0.125	0.5	1	0.5	-	0.25		50.08	0.125	. 	2
Staphylococcus aureus SA1199	0.25	0.5	0.5	2		0.5	2	\$0.06	<u>_</u>	2	0.5
Staphy lococcus aureus SA1199A	80	20.06	S0.06	\$0.06	\$0.06	≥0.06	0.5	80.06	\$0.06	\$0.06	\$0.06
Staphylococcus aureus SA1199B	0.5	-	-	0.5	_	_	-	0.5	0.5	-	7
Staphylococcus haemolyticus 105		1	1	1	1		7	7	' (7)	: 	. 2
Staphylococcus haemolyticus 415	16	7	-	2	7	7	7	. 71	: (3)	. 	2
Staphylococcus epidermidis 270	-	0.5		-	7	-		0.5			
Entercoccus faecium 180		≥0.06	≥0.06	≥0.06	0.125	.12	0.25	0.5	0.125	≥0.06	0.25
Entercoccus faecium 180-1	0	≥0.06	≥0.06	•	0	0.0	\$0.06	90.0≥	0	\$0.05	0
Entercoccus faecalis 2041	≥0.06	≥0.06	≥0.06	20.06	≥0.06	0	0.125	Ö	12	20.06	50.06
Entercoccus faecalis 276		0.125	≥0.06	≥0.06	≥0.06	0.0	2	20.06	0.125	0.25	0.125
Entercoccus gallinarum 245	0.25	7	-	2	≥0.06		7		7	_	. 7
Haemophilus influenzae RD							>64			•	
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	50.06	VI	≥0.06	≥0.06	≥0.06	≥0.06	≥0.0€	50.06	S0.06	≥0.05	50.06
Streptococcus pneumoniae Pl	50.08	S0.06	≥0.06	≥0.06	≥0.06	≥0.06	50.06	≥0.06	30.05	\$0.06	20.06

0.25

0.25 0.5 0.5 0.5 0.25 0.25

0.25 0.25 50.06

0.25

\$0.06

\$0.06

\$0.06

≥0.06 ≥0.06

≥0.06

\$0.06

0.25

≥0.06

>64

>64 ≥0.06

>64 S0.06 <0.06

>64 \$0.06 \$0.06

Streptococcus pyogenes C203 Streptococcus pneumoniae Pl

Escherichia coli EC14

Entercoccus gallinarum 245 Haemophilus influenzae RD

>64

>64

In Vitro Activity of Formula I Compounds MIC (mcg/ml)/Compound TABLE 3 40 45

5

10

15

20

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35

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4											<u>_</u>	_		-					-
104	20.06	≥0.06	0.25	≥0.06	≥0.06	≥0.06	≥0.06	7	≥0.06	0.125	7	7	0.25	\$0.06	≥0.06	20.06	\$0.06	7	. ;
103	0.5	≥0.06		\$0.06	0.5	-4		1	S0.06	7	4	&	1	; ;	\$0.06	\$0.06	0.25	80	
102	-	≥0.06	1	≥0.06	≥0.06	-		0.125	\$0.06		2	7	1		≥0.06	\$0.06	0.25	4	
101	0.5	≥0.06	-	≥0.06	0.25	0.5	0.25	0.25	0.5	-	-4	8	0.5	i	₹0.0€	≥0.06	0.125	a o	!!!
100	-	0.5	0.5			≥0.06		0.5	0.125	0.5		7	0.5	1	20.06	≥0.06	0.25	7	
66	0.5	0.25	2	0.125	0.5	1	0.5	0.5	0.5	0.5	æ	32	1		\$0.06	0.25	0.25	32	
98	5.0	50.05	0.25	≥0.06	20.06	0.25	0.5	0.5	50.06	-	П		≥0.06		\$0.06	50.05	0.125		
97	1	0.25	1	1	0.25	0.5	>64	2	20.06	1	2	2	1		S0.06	\$0.08	≥0.06	2	
96	1	1	1	2	1	1	1	2	20.06	1	2	7	2	0.5	0.25	1	0.5	7	
95	5.0	2	0.5	1		. ~	7	0.5	20.06	1	1		1	0.5	≥0.06	≥0.06	0.125	-	:
Organism	Staphylococcus aureus 446			SI	S	St 'ylococcus aureus 491	Staphylococcus aureus S13E	Staphylococcus aureus SA1199	Staphylococcus aureus SA1199A	Staphylococcus aureus SA1199B	Staphylococcus haemolyticus 105	Staphylococcus haemolyticus 415	27	Entercoccus faecium 180		Entercoccus faecalis 2041	Entercoccus faecalis 276	Entercoccus gallinarum 245	

Organism	106	107	108	109	110	111	112	113	114	115	116
Staphylococcus aureus 446	2	2	2	1	0.5	2	2	≥0.0€	0.5	0.125	0.5
Star sylococcus aureus 489	7	-	0.25	≥0.06	-		0.25	0.125		0.125	٦.
Staphylococcus aureus 447	0.25	-	0.5	-1		-	-	0.25	0.5	0.5	
Staphylococcus aureus X400		1	2	\$0.05	-	-	-	0.125	7		_
Staphylococcus aureus X778		0.5	0.125	S0.06	0.5	7	7	1	7		2
Staphy lococcus aureus 491	0.5	1	0.25	0.25	0.25	7	7	0.25	-	0.5	0,5
Staphylococcus aureus S13E	٦,	2		0.25		-	7	\$0.06	7	0.25	. بـــ
Staphylococcus aureus SA1199		1	2	≥0.06	0.25	7	2	7	2	0.125	7
Staphy lococcus au s SA1199A	50.06	20.06	≥0.06	≥0.06	≥0.06	0.5	0.125	≥0.06	≥0.06	\$0.06	\$0.06
Staphylococcus aureus SA1199B	7	2	2	0.5	0.5		0.5	≥0.06	4	0.25	0.5
Staphylococcus haemolyticus 105		2	2		•		7	4	~ :	7	7
Staphylococcus haemolyticus 415	-	7	1	4	7	4	2	1	7	7	4
270	0.25	0.5	0.125	0.25	2		1	0.25		0.5	
	\$0.06	0.125	0.125	0.25	0.25	≥0.06	≥0.06	\$0.06	20.06	-	\$0.06
	20.06	≥0.06	≥0.06	≥0.06	≥0.06	≥0.06	\$0.06	≥0.06	so.06	≥0.06	≥0.06
	0.125	0.5	≥0.06	≥0.06	≥0.06	20.06	\$0.06	20.06	0.25	≥0.06	≥0.06
Entercoccus faecalis 276	0.5	1	0.5	\$0.06	0.5	0.5	0.5	0.25	-1:	0.125	0.25
Entercoccus gallinarum 245		7	50.06	\$0.06	2	43:	71	-	7	7	₹.
Haemophilus influenzae RD	764	794	>64	32	>64	>64	>64	>64	>64	×64	700
Escherichia coli EC14	>64	>64	>64		>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	\$0.06	50.06	20.06		20.06	≥0.06	≥0.06	50.06	≥0.06	\$0.05	50.06
Streptococcus pneumoniae Pl	\$0.06	\$0.06	\$0.08		\$0.06	\$0.06	50.08	\$0.06	S0.06	S0.06	\$0.06

Organism	117	118	119	120	121	122	123	124	125	126	127
Staphylococcus aureus 446	0.5		2	2	2	-	2	4	7	2	
Staphylococcus aureus 489	0.125	0.25	0.5	2	1	≥0.0€	7	0.25	~	0.25	~
Staphylococcus aureus 447	0.5	0.25	7	-	0.5	0.25		•	~	,	~
Staphylococcus aureus X400	\$0.06	0.25	-	0.25	0.125	0	-	-	~	-	7
Staphylococcus aureus X778	0.25	0.5	2	0.125	0.5	0	-	0.5	2	0.5	
Staphylococcus aureus 491	0.5	20.06	50.06	20.06	≥0.05	0	0	12	-	0.25	
Staphylococcus aureus S13E	\$0.06	0.25	0.25	S0.06	٠.	90.0≥	\$0.0¢	≥0.06	-	0.5	. 7
Staphylococcus aureus SA1199		2	2			.0	7		0.5	0.125	7
Staphy lococcus aureus SA1199A		20.06	0.25	\$0.06	20.06	. 0	_		0.25	0.25	0.25
Staphylococcus aureus SA1199B		\$0.08	0.5	0.125	0.25		0.5	20.06	2		~
Staphylococcus haemolyticus 105	•	1	2	2	~	_	1	7	4	0.5	7
Staphylococcus haemolyticus 415	7	-	7	2	7		-		7	7	4
Staphylococcus epidermidis 270	0.5	1	2	2	1	0	-	0.25	, ,		≥0.06
		0.125	0.125	\$0.05	\$0.08	≥0.06		≥0.06	-4	-	\$0.06
Entercoccus faecium 180-1	٠.;	S0.06	≥0.06	≥0.06	20.06	0	≥0.06	\$0.06	<0.06	0	30.05
Fitercoccus faecalis 2041	\$0.06	≥0.06	≥0.06	≥0.06	≥0.06	0		≥0.06	>0.0€	≥0.06	≥0.06
Entercoccus faecalis 276	~	S0.06	0.125	20.06	\$0.06	0	~	20.06		-4	≥0.06
Entercoccus gallinarum 245		1	7	7	7	-:		1	12	7	4
Haemophilus influenzae RD		16	16	16	16	91	16	16			×64
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64			× 64
Streptococcus pyogenes C203	≥0.06	\$0.06	\$0.06	≥0.06	S 0.06	50.06	VI	≥0.06		20.06	20.06
Streptococcus pheumoniae Pl	<0.05	<0.0>	90 0>	<0.05	40 0K	<0.05	90 0×	<0.05	<0 0×	90 08	

Organism	128	129	130	131	132	133	134	135	136	137	138
Staphy lococcus aureus 446	♥.	2	1	2	, party	7	2		\$0.05	0.25	0.125
Staphylococcus aureus 489		≥0.06	0.5		i	7	0.5	0.125	≥0.06	≥0.06	\$0.06
Staphylococcus aureus 447	-1		1	1	7				7		~
Staphylococcus aureus X400		0.25	0.5	1		0.5	0.25	≥0.0€	\$0.06	≥0.06	≤0.0606
Staphylococcus aureus X778		0.25	1	0.5	2	2	!		≥0.06	≥0.06	0.25
Staphylococcus aureus 491	7	0.5	0.5	0.125	0.5	0.25	0.5	0.25	0.25	0.25	0.125
Staphylococcus aureus S13E		0.25	0.5	1		-	2	-	≥0.06	≥0.06	\$0.06
Staphy lococcus aureus SA1199	0.5	0.25		0.25	-	0.25	0.25	-		\$0.06	\$0.05
SA1199	0.5	≥0.06	≥0.06	≥0.06	0.25	0.25	7	\$0.06		50.06	≥0.06
_	7	0.25	7	1	2	7	2	0.25		≥0.06	0.5
S	7	7	-1	1	1	~	2	0.5	7	7	7
s 41	7	4	2	2	7	~~	4	N	4		. 00
7			L	1	2		7	0.5		0.5	7
İ		4	1	20.06	0.25	-	0.5		. ~	0.125	4
Entercoccus faecium 180-1	0.125	≥0.06	S0.06	≥0.06		≥0.06	≥0.06	50.06	20.06	S 0.06	\$0.05
Entercoccus faecalis 2041	N.	\$0.06	_	\$0.05	, ,;	0.25	0.25	12	≥0.05	20.06	≥0.06
Entercoccus faecalis 276	-	0.125	-	0.25	-4		-	0.5	0	\$0.06	\$0.06
Entercoccus gallinarum 245	7	0.125	2	7	~ ~ ;	7	₹	7	4	œ	0.125
Haemophilus influenzae RD		>64 -									
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	\$0.06	S0.06	\$0.06	20.06	50.06	≥0.06	50.06	S0.06	\$0.06	20.06	50.06
Streptococcus pneumoniae P1	50.06	≥0.06	50.08	20.06	≥0.08	≤0.06	S0.06	_	20.06	S0.06	50.06

Organism	139	140	141	142	143	144	145	146	147	148	149
Staphylococcus aureus 446	0.5	0.125	2	2	9.0	16	0.5	9.0	0.5	1	0.5
Staphylococcus aureus 489	0.25	≥0.06	0.25	0.5	\$0.06	4	≥0.06	0.25		0.25	≥0.06
Staphylococcus aureus 447	-	0.25	1	2	4	16	-	2	0.125	1	4
Staphy lococcus aureus X400	0.25	≥0.06	0.25	1	0.125	œ	0.25	0.5	4	≥0.06	≥0.06
Staphylococcus aureus X778	0.125	0.25	0.5	-	20.06	60	0.125	50.06	0.25	2	0.5
Staphylococcus aureus 491	0.5	0.25	0.5	0.5	0.5	æ	٥. د	1	20.06	0.125	0.5
_E	90.05	≥0.06	0.25	2	0.125	œ	0.125	0.5		1	0.25
Staphylococcus aureus SA1199	0.125	20.06	0.25	1	0.125	80	0.25	50.06	0.5	2	0.25
Staphy lococcus aureus SA1199A	\$0.08	≥0.06	20.06	≥0.06	20.06	7	≥0.06	20.06	0.25	\$0.06	\$0.06
Staphylococcus aureus SA1199B	2	S0.06	2	2	0.25	80	\$0.06	≥0.06	≥0.06	0.5	
Staphylococcus haemolyticus 105	4	2	1	1	80	64	7	7	-	Н	7
Staphylococcus haemolyticus 415	œ	8	Þ	1	32	>64	æ	4	æ	7	16
Stringlococcus epidermidis 270		0.25	1	0.25	7	16		2	16	0.5	-
	2	1	5.0	5.0	4	ω	4	œ	7	0.25	
Entercoccus faecium 180-1	\$ 0.06	≥0.06	≥0.06	VI	≥0.06	4	≥0.06	20.06	7	≥0.06	90 us
Entercoccus faecalis 2041	≥0.06	≥0.06		\$0.06	0.125	80	0.25	0.5	\$0.06	S0.06	20.06
Entercoccus faecalis 276	-	0.5	0.5	1	0.25	œ·	0.125	1	0.125	\$0.06	\$0.06
Entercoccus gallinarum 245	00	© 1	7	7	32	4	0.25	0.5	0.125	7	16
Haemophilus influenzae RD	!					i	1	>64			
	>64	_	>64	>64	>64	>64	>64	>64	>64	>64	> 64
C203	\$0.08	VI	≥0.05	≥0.06	yu'05	0.5	≥0.06	≥0.06	S0.06	≥0.06	\$0.06
	SO 06	_	<0.05	<0.05	V	<0.05	<0.05	S0.06	20.06	<0.05	<0.05

TABLE 3

In Vitro Activity of Formula I Compounds MIC (mcg/ml)/Compound

Organism	150	151	152	153	154	155	156	157	158	159	160
Staphylococcus aureus 446	1	2	2	0.5	2	2	2	0.5	2	0.5	2
Staphylococcus aureus 489	0.5	≥0.06	0.5	1		0.5	1	0.5	2	0.25	0.21
Staphylococcus aureus 447	0.5	-	8	0.5	7	ھ	1	0.25	4	-7	-
Staphylococcus aureus X400	≥0.06	≥0.06	1	0.5	2	1	2	0.5	4	4	₹
	7	1	0.5	0.5	0.5	≥0.06	-	0.25	4	2	4
Staphylococcus aureus 491	\$0.06	0.5	1	0.125	0.5	1	1	≥0.06	1	2	0.125
Staphylococcus aureus S13E	0.25	0.25	0.5	0.125	0.25	-	1	0.25	2	1	
Staphylococcus aureus SA1199	П	0.125	1	0.5	2	-	1	1	7	0.125	0.25
Staphylococcus aureus SA1199A	≥0.06	≥0.05	0.25	≥0.05	0.125	≥0.06	\$0.06	≥0.06	-	20.06	0.125
Staphylococcus aureus SA1199B	0.5	0.25	0.5	0.25	0.25		0.5	1	4	20.06	20.06
Staphylococcus haemolyticus 105		-1	16	7	7	16	4	П	4	16	
Staphylococcus haemolyticus 415	7	4	16	1	7	16	2	1	~	6 0	80
Staphylococcus epidermidis 270	0.25	0.5	4	0.25	0.5	1	1	0.25	4	0.5	-
Entercoccus faecium 180	0.25	0.25	4	0.125	1	4	1	S0.06	0.25	~1	0.5
Entercoccus faecium 180-1	20.06	≥0.06	0.125	S0.06	20.06	≥0.0€	\$0.06	≥0.06	0.25	≥0.06	≥0.06
Ent reoccus faecalis 2041	20.06	≥0.06	0.125	90.05	0.125	0.125	0.5			0.125	\$0.05
Entercoccus faecalis 276	-	≥0.06	0.25	0.5	0.5	0.25	2	80.06	7	0.125	7
Entercoccus gallinarum 245	7	4	16		9	16	7	-	60	· œ	: 00
Haemophilus influenzae RD						16	7			•	
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	\$0.06	20.06	≥0.06	≥0.06	≥0.06	≥0.06	50.05	≥0.06	≥0.06	\$0.06	50.08
Streptococcus pneumoniae P1	20.06	≤0,06	≥0.06	≥0.06	≥0.06	≥0.06	≥0.06	50.05	50.08	≥0.06	\$0.06

Organism	161	162	163	164	165	166	167	168	169	170	171
Staphylococcus aureus 446	0.5	0.5	1	2	1	7	1	≥0.06	0.25	2	7
Staphy lococcus aureus 489	\$0.05	0.25	æ	7	7	7	16	0.125	≥0.06	0.25	0.5
Staphylococcus aureus 447	-	≥0.06	0.5	2	0.5	7	7	20.06	2	0.5	
Staphylococcus aureus X400	0.5	≥0.06	5.0	0.5	0.5	,	1	≥0.06	≥0.06	0.5	
Staphylococcus aureus X778	•	≥0.06	2	1	0.125	1	16	0.5	50.06	1	
Staphylococcus aureus 491	0.5	0.25	S0.06	1	0.5	0.5	2	0.5	0.25	0.5	0.25
Staphylococcus aureus S13E	0.125	≥0.06	1	4	≥0.06	4	4	≥0.06	≥0.06	0.25	\$0.08
Staphylococcus aureus SA1199	0.25	50.06	2	2	0.25	7	2	0.5	≥0.06	. —	0.25
Staphylococcus aureus SA1199A	≤0.06	≥0.06	0.5	٠.٥	50.05	0.125	4	≥0.0€	≥0.06	≥0.06	\$0.06
Staphylococcus aureus SA1199B	0.25	\$0.08	1	2		2	4		0.125	0.25	0.25
Staphylococcus haemolyticus 105	4	0.25	8	2	4	2	32	0.5	7	4	7
Staphylococcus haemolyticus 415	80	2	8	2	7	2	16	2	4	4	æ
Staphylococcus epidermidis 270	1	≤0.06	4	1	1	0.5	8	0.125	0.25	1	-
Entercoccus faecium 180	2	≥0.05	1	0.5	0.5	0.25	2	0.25	1	2	
Entercoccus faecium 180-1	\$0.06	50.06	S0.06	S0.06	≥0.0€	20.06	≥0.0€	≥0.0€	S0.06	≥0.06	0
Entercoccus faecalis 2041	50.06	\$0.06	1	1	≥0.06	50.06	80	50.08	≥0.06	0	SO.06
Entercoccus faecalis 276	0.125	\$0.08	1	1	0.5	S	4	0.125	≥0.06	0.5	12
Entercoccus gallinarum 245	80	2	80	2	7	7	16	2	4	4	i co
Haemophilus influenzae RD										>64	>64
Escherichia coll EC14	>64	>64	>64	>64	>64	>64 4	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	\$0.06	50.06	20.06	≥0.06	≥0.06	≥0.06	0.25	S0.06	\$0.08	S0.08	≥0.06
Strentococcus pheumoniae Pl	90 0×	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	90 0V	90 00	×0 06	×0 0×	A0 05	40 0X	20 05	20 05	20 05

Staphylococcus aureus 446 4.5 Staphylococcus aureus 489 0.5 Staphylococcus aureus 447 0.5	V						-		+0+	701
	7	0.5		2	0.5	П	0.125	0.125	50.05	2
	2	≥0.06	0.25	0.5	≥0.06	0.125	≥0.06	≥0.06	≥0.06	7
	4	\$	-1		4	0.5	0.25		• •	0.25
Staphylococcus aureus X400 0.5	7	≥0.06	0.125	-	≥0.06	0.125	≥0.06	≥0.06	0	
Staphylococcus aureus X778 2	•	20.06	0.5	-	7		\$0.06	0.	0	7
Staphylococcus aureus 491 0.5	7			7	0.5	0.125	12	v		
S0.0	6 4	80.05	0.25	2		0.5	0.25	≥0.06	≥0.06	0.25
1 1	7	\$0.06	≥0.06	2	0.25	1	1	0.125	≥0.06	7
A 50.0	6 0.5	30.05	0.5	>64	-	≥0.06	0.0	≥0.06	\$0.06	50.06
B 50.0	9	0.125	\$0.05	-		7	0.125	50.06		7
105 0.2	5 2	4	2	4	4	1	0.5	2	0.25	7
415 2	4	16	4	7	16	2	1	7		4
Staphylococcus epidermidis 270 0.5	2	2	0.5	0.5	1	0.25	0.25	0.125	0.125	0.25
Entercoccus faecium 180 0.5	0.5	2	1	7	4	0.25	20.06	8	4	2
-1 50.		≥0.06	\$0.06	≥0.06	≥0.06	≥0.06		0.125	≥0.06	≥0.06
\$0.	9:	≥0.06	\$0.06	0.125			• •	0.25	2	
Entercoccus faecalis 276 0.12		20.06	\$0.08	7	0.25		0.0		~	,
5	3	16	Þ	~	~ ;	7		0.25	\$0.06	-
Haemophilus influenzae RD 32	>64	>64	16	æ	>64	4	7		32	*9
Escherichia coli EC14 >64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203 50.06	90.05	≥0.06	0.5	0.25	16	\$0.05	\$0.06	\$0.06	≤0.06	20.06
Streptococcus pneumoniae P1 50.0	90.05 90	50.06	0.5	0.25	9	S0.06	\$0.06	≥0.06	≥0.06	\$0.06

Organism	183	184	185	186	189	190	161	192	193	194	195
Starthylococcus aureus 446	≥0.06	2	≥.06	≥.06	0.5	0.25	2	0.5	9.0	≥0.06	0.5
Staphylococcus aureus 489	≥0.06	≥.06	₹.06	≥.06	-	0.125	2	1		0.125	
Staphylococcus aureus 447	≥0.06	≥.06	≥.06	2.06	0.5		2	2	0.0	0.5	
Staphylococcus aureus X400	≥0.06	0.5	≥.06	≥.06	0.125	≥0.06	1	1	0.25	20.06	7
Staphylococcus aureus X778		• :	2.06	5.06	0.25	_	2	→	0.	Ö	0.5
staphylococcus aureus 491	0.125	0.5	≥.06	3.06	>0.05	0.125	1	0.5	\$0.06	\$0.06	0.5
Staphylococcus aureus S13E	50.06	1	≥.06	S.06	0.5	0.125	7	1	\$0.06	0	· 10
Staphylococcus aureus SA1199	\$0.08	0.125	≥.06	S.06	0.5	0.25	2	7	0.125	0.5	0.5
Staphylccoccus aureus SA1199A	\$0.06	≥.06	≥.06	이		≥0.06	0.5	0.75	0	20.06	. •
Staphylococcus aureus SA1199B	\$0.06	1	≥.06	≥.06	1	2	2		27	0.125	-
Staphylococcus haemolyticus 105	\$0.06	0.25	S.06	0.5		∞.	7	-1	0.5	1	
Staphylococcus haemolyticus 415	\$0.06	0.	≥.06	-	-	.	80	7	-	7	. 4
Staphylococcus epidermidis 270	\$0.00	4	≥.06	0.125	0.25	2	2		0.25	0.5	0.25
Entercoccus i eclum 180	2	80	0.125	7	0.125	æ	4	0.25	\$0.06	0	0.5
Entercoccus faecium 180-1	\$0.06	≥.06	S .06	≥.06	≥0.06	0	0.25	. 12	0	≥0.06	≥0.06
Entercoccus faecalis 2041	\$0.06	1	5.06	s.06			1	0.125	20.06	0	20.06
Entercoccus faecalis 276	0.125	0.5	≥.06	≥.06	0.25	12	4	0.5		12	0.25
Entercoccus gallinarum 245	0.5	4	2.06	7	1	æ	œ	7	1	2	47
Haemophilus influenzae RD	>64	64	8		32	۰	>64	>64	>64	>64	32
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203	\$0.06		2.06	≥.06	≥0.06	20.06	50.06	\$0.06	20.06	\$0.06	≥0.06
Streptococcus pneumoniae P1	\$0.06		≥.06	S .06	\$0.06	S0.06	50.06	50.05	\$0.05	80.06	≥0.06

TABLE 3
In Vitro Activity of Formula I Compounds
MIC (mcg/ml)/Compound

			272	7/7	222	404	202	203	204	205
Staphylococcus aureus 446 0	5.	1	1	0.5	1	7	4	0.5	0.125	7
Staphylococcus aureus 489		2	0.125	2	0.25	8	7	0.5	0.25	0.5
Staphylococcus aureus 447 0	.5	7	0.125	1	0.5	16	æ	-	≥0.06	0.5
Staphylococcus aureus X400 0	.5	7	0.5	2	-	4	4		0.125	0.5
Staphylococcus aureus X778	-	2	0.125	1	0.5	4	4	-	0.5	0.5
Staphylococcus aureus 491 0.	.25	1	>0.06	0.5	0.125	7	œ		≥0.05	0.5
Staphylococcus aureus S13E	-	2	0.125	0.5	0.5	8	4	2	0.5	0.5
Staphylococcus aureus SA1199 0	5.0	2	0.5	1		æ	80	2	0.125	,
9.A	SO.06	7	30.05	0.125	20.06	2	7	0.5		\$0.08
Staphylococcus aureus SA1199B 0	5.5	2	0.5	1	-	16	œ		0.25	0.5
Staphylococcus haemolyticus 105 0	5.0	1	0.5	2	-	80	7	~	0.5	-
Staphylococcus haemolyticus 415	-	ð	1	7	7	œ	œ	7	0.25	
Staphylococcus epidermidis 270 0	0.25	0.5	0.25	0.5	0.25	7	4	0.5	\$0.06	0.125
Entercoccus faecium 180	5.5	0.5	20.06	0.5	0.25	0.5	0.5	0.125	0.25	0
Entercoccus faecium 180-1	0.06	0.25	≥0.06	≥0.06	\$0.06	0.5	0.5	20.06	0.125	\$0.06
	90.0	0.25	≥0.0€	\$0.06	\$0.06	~		0.25	\$0.06	0.25
Entercoccus faecalis 276 0	25	7	0.25	-	0.5	4	4	5.0	S0.06	0.5
Entercoccus gallinarum 245	-	•		4	~	œ.	60	~	0.25	_
Haemophilus influenzae RD	32	32	32	32	32	32	32	16	2	16
	>64	>64	>64	>64	>64	>64	>64	>64	>64	794
Streptococcus pyogenes C203 S0	50.06	≥0.06	50.05	S0.06	50.06	\$0.06	\$0.06	\$0.06	50.06	\$0.06
Streptococcus pneumoniae Pl S0	90.	≥0.06	≥0.06	≥0.06	≥0.0€	\$0.08	≥0.06	S0.06	≤0.06	\$0.06

TABLE 3 In Vitro Activity of Formula I Compounds

•	nd	
	/Compoun	
	(mcg/m])/(
	MIC	

Organism	206	207	208	209	210	211	212	213	214	215
Staphylococcus aureus 446	0.5	8	1	1	2		S0.06	≥0.06	-	0.5
Staphylococcus aureus 489	~	7	0.5	1	٦	0.25	≥0.06	≥0.06		2
Staphylococcus aureus 447	0.5	æ	-	1	0.5	0.5	0.25	0.25	2	0.5
Staphylococcus aureus X400	0.5	8	0.25	1	≥0.06	0.5	≥0.06	≥0.06	0.5	0.5
Staphylococcus aureus X778	0.5	80	0.125	1	1	1	≥0.06	≥0.06	-	≥0.06
Staphylococcus aureus 491	\$0.06	1	0.5	0.25	\$0.06	0.25	\$0.06	\$0.06	-	0.25
Staphylococcus aureus S13E	-1	æ	0.25	0.5	≥0.06	0.5	20.06	\$0.06		7
Staphylococcus aureus SA1199	0.5	æ	0.5	0.25	0.5	0.5	20.06	≥0.06	0.5	≥0.06
Staphylococcus aureus SA1199A	≥0.06	4	S0.06	≥0.06	≥0.06	0.125	20.06	\$0.08	0.5	0.5
Staphylococcus aureus SA1199B		16	0.5	0.5	0.125	-	\$0.06	20.06		
Staphylococcus haemolyticus 105	0.5	8	0.25	0.5	-	0.5	-	0.5	-	7
Staphylococcus haemolyticus 415	1	-	2	-	1	0.5	-	2	7	
Staphylococcus epidermidis 270	0.25	8	0.5	0.125	0.25	0.5	\$0.06	0.5	90.0₹	0.125
Entercoccus faecium 180	≥0.06		0.25	≥0.06	\$0.06	\$0.06	20.06	0.125	0.25	20.06
Entercoccus faecium 180-1	≥0.06	≥0.06	\$0.06	\$0.06	≥0.06	\$0.06	≥0.06	\$0.06	≥0.0€	\$0.06
Entercoccus faecalis 2041	0.25	0.125	\$0.06	\$0.06	\$0.06	50.06	\$0.06	\$0.06	0.125	0.25
Entercoccus faecalis 276	20.06	0.25	0.125	0.25	\$0.06	\$0.06	\$0.06	\$0.06	0.25	7
Entercoccus gallinarum 245	****	-	2	1	1	\$0.06		7	2	64
Haemophilus influenzae RD	1		32	16	>64	>64	>64	32	32	>64
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Streptococcus pyogenes C203			≥0.06	\$0.08	\$0.06	\$0.06				S 0.06
Streptococcus pneumoniae Pl	\$0.06	S0.06	S0.06	\$0.05	\$0.06	\$0.06	20.06	50.06	\$0.06	\$0.06

Organism	216	217	218	219	220	221	222	223	224	225
Staphylococcus aureus 446	1	0.25	Þ	8	1	1		0.25	0.5	-
Staphylococcus aureus 489	1	≥0.05	1	8	0.5	0.25	0.125	1	0.25	2
Staphylococcus aureus 447	1	1	1	8	0.5	0.5	0.5	0.5	0.5	1
Staphylococcus aureus X400	1	S0.06	0.25	8	0.5	0.5	0.125	-	0.125	-
Staphylococcus aureus X778	0.25	≥0.06	1	8	0.5	0.5	\$0.06	1	0.125	0.5
Staphylococcus aureus 491	1	0.25	0.5	4	20.06	0.125	0.125	0.125	0.125	7
Staphylococcus aureus S13E	-	\$0.08	32	8	0.5	0.5	50.06	0.5	0.25	7
Staphylococcus aureus SA1199	≥0.06	≥0.06	Þ	4	-	-	1	2	0.25	1
Staphylococcus aureus SA1199A	1	S0.06	90.05	1	≥0.06	>0.05	0.125	\$0.06	≥0.06	0.25
	0.5	0.125	0.25	8	0.5	1	0.125	1	0.5	7
Staphylococcus haemolyticus 105	5.0	2	0.5	2	0.5	1	1	1	1	0.5
Staphylococcus haemolyticus 415	0.25	8	Þ	2	9.0	2	1	1	0.5	4
Staphylococcus epidermidis 270	0.125	0.5	1	4	1	0.125	0.5	0.5	0.25	1
Entercoccus faecium 180	≥0.0€	2	\$0.06	1	0.125	50.06	S 0.06	≥0.06	≥0.05	≥0.06
Entercoccus faecium 180-1	≥0.06	\$0.08	≥0.06		≥0.06	≥0.06	S0.08	50.08	S0.06	≥0.06
Ent reoccus faecalis 2041	0.25	\$0.06	0.25	2	≥0.06	≥0.06	S0.06	≥0.06	S0.06	0.125
Entercoccus faecalis 276	0.5	\$0.05	≥0.06	2	0.125	0.25	≥0.06	0.125	≥0.06	0.25
Ent reoccus gallinarum 245	64	80	\$0.06	2	0.5	7		-	0.5	₹.
Haemophilus influenzae RD	>64	>64	>64	32	>64	32	32	>64	32	× 64
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	764	× 1
Streptococcus pyogenes C203	≥0.06	S0.06	\$0.06	\$0.06	\$0.06	20.06	≥0.06	≥0.06	\$0.08	50.06
Streptococcus pneumoniae Pl	\$0.06	20.06	20.06	20.06	≥0.06	\$0.06	20.06	\$0.06	\$0.06	\$0.06

Organism	226	227	228	229	230	231	232	233	234	235
Staphylococcus aureus 446	1	2	7	1	0.25	0.25	4	4	4	0.5
Staphylococcus aureus 489	0,5	2	2	1	0.25	\$0.08	æ	4	7	0.5
Staphylococcus aureus 447	0.5	2	4	2	0.5	0.25	16	16	: œ	0.25
Staphylococcus aureus X400	C)	1	1	1	0.5	≥0.05	æ	œ	œ	0.125
Staphylococcus aureus X778	0.25	Ą	Þ	1	0.25	S0.06	œ	œ	♂	0.5
Staphylococcus aureus 491	(1)	2	1	0.5	0.125	\$0.06	4	œ	00	0.125
Staphylococcus aureus S13E	0.5	4	8	,4	0.5	≥0.06	œ	8	&	0.125
Staphylococcus aureus SA1199	-4	4	Þ	1	0.25	≥0.05	16	32	60	0.25
Staphylococcus aureus SA1199A	0.125	9.0	\$0.05	≥0.06	≥0.06	≥0.05	2	4	~	≥0.06
Staphylococcus aureus SA1199B		4	4		0.25	\$0.05	32	16	æ	0.5
Staphylococcus haemolyticus 105	7	2	2	٦	1	\$0.05	2	>64	80	0.5
Staphylococcus haemolyticus 415		4	4	2	2	0.5	32	>64	16	7
Staphylococcus epidermidis 270		2	2	0.5	0.5	0.125	60	80	7	0.5
Entercoccus faecium 180	20.06	0.25	1	≥0.06	S0.06	50.06	0.5	7	-	0.5
Entercoccus faecium 180-1	\$0.06	\$0.05	≥0.05	≥0.06	\$0.06	50.06	1	7	-	\$0.06
Entercoccus faecalis 2041	80.08	0.25	0.25	≥0.06	\$ 0.0 6	50.06	2		0.5	90.05
Entercoccus faecalis 276	0.25	0.5	1	0.25	≥0.06	≥0.06	8	æ	7	0.125
Entercoccus gallinarum 245	1	4	4	2	2	0.5	32	>64	16	
Haemophilus influenzae RD	32	>64	>64	2	32	32	16	>64	>64	- α
Escherichia coli EC14	>64	>64	>64	>64	>64	>64	>64	>64	>64	> 64
Streptococcus pyogenes C203	\$0.06	\$0.06	0.125	\$0.0 8	50.06	50.06	≥0.06			20.06
	\$0.06	\$0.06	\$0.05	90.0≥	≥0.05	90.0≥	≥0.06	0.5	0.25	\$0.06

Organism	236	237	238	239	240	241
Staphylococcus aureus 446	1	2	-			0.5
Staphylococcus aureus 489	4	0.5	0.5	0.5	-1	0.5
Staphylococcus aureus 447	7	7	0.5	0.5	0.5	1
Staphylococcus aureus X400	2	1	-	0.25	0.25	0.5
aureus	2	0.5	0.5	0.25	0.5	1
	4	0.25	0.25	0.25	0.25	0.25
Staphylococcus aureus S13E	Þ	0.25	0.125	0.5	0.5	0.25
aureus	4	1	0.5	0.5	5.0	-
Staphylococcus aureus SA1199A	2	≥0.06	\$0.06	50.06	50.06	50.06
	4	0.25	0.5	0.5	0.25	-
Staphylococcus haemolyticus 105	4	1	0.5	1	1	1
Staphy lococcus haemolyticus 415	4	1	2	1	2	1
Staphylococcus epidermidis 270	2	0.5	0.5	0.25	52.0	0.5
Entercoccus faecium 180	1	0.25	0.125	≥0.06	≥0.05	≥0.06
Entercoccus faecium 180-1	1	\$0.06	\$0.06	≥0.06	90.0≥	\$0.08
Entercoccus faecalis 2041	1	0.125	20.06	≥0.06	20.06	50.05
Entercoccus faecalis 276	2		≥0.06	0.25	0.5	50.05
Entercoccus gallinarum 245	4	-	\$0.06	1	7	50.05
Haemophilus influenzae RD	32	æ	794	>64	>64	>64
Escherichia coli EC14	>64	>64	>64	>64	>64	×64
Streptococcus pyogenes C203	S0.06	\$0.06	\$0.06			50.06
Streptococcus pheumoniae Pl	20.06	80.08	\$0.08	20.06	\$0.08	50.06

The formula \underline{I} compounds have also shown $\underline{in \ vivo}$ antimicrobial activity against experimentally-induced infections in laboratory animals. When two doses of test compound were administered to mice experimentally infected with the test organism, the activity observed was measured as an ED₅₀ value (effective dose in mg/kg to protect 50% of the test animals: see W. Wick $\underline{et} \ \underline{al}$., \underline{J} . Bacteriol. 81, 233-235 (1961)). ED₅₀ values observed for illustrative compounds are given in Table 4.

TABLE 4

5	In Vivo	Activity of F (mg.	ormula I Compo /kg/2)	unds ED50
·			Streptococcus	Streptococcus
	Compound	aureus	pyogenes	pneumoniae
	vancomycin	1.2	0.8	1.1
	A82846A	0.19	0.084	0.39
10	A82846B	0.25	0.12	0.18
	A82846C	1.3	1.5	4.6
	1	0.086	0.052	0.025
	2	0.27	0.014	0.025
	4	0.36	0.012	0.036
15	5 .	0.13	0.039	0.036
	6	0.15	0.013	0.021
	8	0.12	>0.5	0.273
	12	0.13	>0.5	>0.5
20	14	0.43	0.37	>0.5
20	22	0.049	>0.5	>.05
	25	0.16	0.087	0.088
	29	0.088	0.1	0.054
	32	0.055	0.034	0.039
25	36	0.19	0.28	0.31
	39	0.1	0.045	<0.031
	41	n.d.	0.082	0.087
	46	n.d.	0.378	0.156
	49	0.053	0.045	<0.031
30	50	0.1	0.047	0.057
	51	0.16	0.057	0.036
	52	0.052	0.046	0.074
	53	0.077	0.16	0.071
35	57	0.041	0.054	0.046
30	64	n.d.	0.044	<0.031
	87	n.d.	0.054	0.027
	90	n.d.	0.058	0.049
	93	n.d.	0.074	0.012
40	94	n.d.	0.16	0.049
	97	n.d.	0.066	0.038
	100	n.d.	0.062	0.046
	104	n.d.	0.12	0.041
	105	n.d.	0.12	0.041
45	106	n.d.	0.2	0.036
	107	n.d.	0.27	0.092
	108	n.d.	0.046	0.041
	111	n.d.	0.099	0.084
50	114	n.d.	0.091	0.76
•••	116	n.d.	0.89	0.058
	118	n.d.	0.91	0.046
	119	n.d.	0.16	0.08
	120	n.d.	0.058	0.005
55	121	n.d.	0.041	0.047

TABLE 4

In Vivo Activity of Formula I Compounds ED50 (mg/kg/2)Stapy lococcus Streptococcus Streptococcus Compound pneumoniae pyogenes aureus 0.31 122 n.d. 0.23 0.039 0.076 123 n.d. 0.092 0.041 n.d. 124 0.077 <0.031 131 n.d. 0.046 <0.031 n.d. 204 <0.031 0.041 211 n.d. <0.031 223 n.d. <0.031 0.078 229 n.d. 0.058 0.046 0.078 n.d. 230 n.d. = not; done

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One important aspect of the antimicrobial activity of many of the formula I compounds is their activity against vancomycin-resistant enterococci. This activity is illustrated in Table 5, which summarizes a comparison of the activity of illustrative compounds against representative vancomycin-resistant and vancomycin-susceptible enterococci (*Enterococcus faecium* and *Enterococcus faecalis*, mean geometric MIC (mcg/mL)), as determined using the standard broth micro-dilution assay. End points were read after 24-hour incubation. Modification of the amino sugar of the disaccharide moiety provides improved activity against vancomycin-resistant strains over the parent glycopeptide antibiotic.

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45

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TABLE 5

Compound No.	Vanc mycin Resistant Strains	Vancomycin Sensitive Strains
vancomycin	282	3.9
A82846A	>64	1.7
A82846B	29	0.22
A82846C	353	1.3
1	0.25	0.0061
2	0.044	0.00038
3	2.8	0.11
4	0.50	0.062
5	0.50	0.072
6	1.2	0.14
7	2.8	0.43
8	1.0	0.57
9	11	0.38
10	3.4	3.5
11	6.7	0.22
12	1.7	1.1
13	19	0.76
14	0.50	0.76
15	6.7	0.14
16	9.5	0.67
17	9.5	0.38
18	6.7	0.38
19	4.8	0.22
20	4.8	0.38
21	5.7	4.3
22	1.0	1.5
23	5.7	2.0
24	54	0.67
25	4.0	0.22
26	54	0.66
27	45	1.5
28	4.7	0.71
29	0.21	0.031
30	4.7	0.071
31	9.5	1.2
32	0.50	0.089
33	2.8	0.18
		3.4
34	4.0	0.25
35	5.6	
36	0.25	0.21
37	2.4	0.25
38	4.0	0.42
39	1.2	0.31
40	0.50	0.31

55

0.84

0.21

TABLE 5

5	Compound No.	Vancomycin Resistant Strains	Vancomycin Sensitive Strains
	42	1.7	0.089
10	43	13	1.1
	44	13	0.50
	45	2.0	0.50
	46	0.71	0.50
	47	4.7	0.57
15	48	4.8	0.50
	49	0.71	0.083
	50	0.12	0.054
	51	0.84	0.22
20	52	0.59	0.11
20	53	0.35	0.25
	54	1.7	0.56
	55	13	1.7
	56	19	1.0
25	57	0.35	0.041
	58	5.7	0.76
	59	51	0.42
	60	19	3.0
	61	16	0.65
30	62	9.5	0.22
	63	54	0.66
	64	0.71	0.077
	65	2.4	0.20
35	66	16	0.76
35	67	1.7	0.16
	68	6.7	0.25
	69	13	0.44
	70	2.0	0.092
40	71	11	0.57
	72	4.7	0.28
	73	11	0.25
	74	11	0.50
	75	16	0.30
45	76	8.0	0.76
	78	16	0.042
	79	0.84	0.042
	80	1.7	0.23
E 0	81	1.0	0.50
50	82	22	1.7
	83	54	0.66
	84	23 3.4	0.11
	85	1.4	0.036
55	86	0.71	0.047
	87	J . / 1	I

TABLE 5

5	Compound No.	Vancomycin Resistant Strains	Vancomycin Sensitive Strains
	88	1.7	0.055
10	89	11	0.44
10	90	0.71	0.041
	91	2.8	0.11
	92	1.7	0.082
	93	0.42	0.042
15	94	0.50	0.041
	95	1.7	0.054
	96	1.4	0.11
	97	0.71	0.054
	98	2.4	0.095
20	99	72	0.76
	100	0.71	0.042
	101	4.0	0.25
	102	2.0	0.13
	103	4.0	0.33
25	104	1.2	0.062
	105	0.84	0.062
	106	0.71	0.034
	107	0.59	0.082
30	108	0.84	0.04
	109	72	0.22
	110	1.7	0.047
	111	0.71	0.031
	112	1.4	0.072
35	113	0.84	0.054
	114	0.59	0.031
	115	8.0	0.19
	116	0.42	0.031
	117	4.8	0.14
40	118	0.84	0.048
	119	0.59	0.048
	120	1.0	0.072
	121	1.0	0.063
45	122	1.0	0.054
	123	1.0	0.041
	124	0.84	0.047
	125	3.4	0.14
	126	2.4	0.11
50	127	1.2	0.33
	128	2.0	0.11
	129	27	1.52
	130	4.8	0.22
	131	0.84	C.028
55	132	1.2	0.048

TABLE 5

5	Compound No.	Vancomycin Resistant Strains	Vancomycin Sensitive Strains
	133	4.0	0.13
	134	2.0	0.13
10	135	4.8	0.22
	136	23	0.76
	137	6.7	0.38
	138	38	0.87
	139	23	0.38
15	140	6.7	0.19
	141	8.0	0.25
	142	45	1.5
	143	2.0	0.048 .
20	144	11	9.2
20	145	64	1.3
	146	64	1.5
	147	25	1.3
	148	0.15	0.052
25	149	45	0.66
	150	1.7	0.25
	151	4.5	0.14
	152	27	1.2
	153	1.4	0.083
30	154	2.8	0.072
	155	128	1.3
	156	5.7	0.17
	157	2.0	0.054
35	158	1.7	1.0
	159	27	0.50
	160	9.5	0.22
	161	23	0.44
	162	4.8	0.12
40	163	2.0	0.11
	164	1.7	0.062
	165	4.0	0.055
	166	1.7	0.055
	167	1.0	0.10
45	168 169	3.4	0.50
	170	8.0	0.22
	171	9.5	0.22
	172	3.4	0.13
50	173	2.0	0.12
	174	19	0.76
	175	9.5	0.22
	175	1.2	1.13
	178	2.8	0.13

TABLE 5

5	Compound No.	Vancomycin Resistant Strains	Vancomycin Sensitive Strains
	179	1.7	0.060
10	180	>128	0.71
	181	8.0	0.060
	182	13	0.250
	183	23	0.130
	184	27	0.570
15	185	4.7	0.060
	186	11	0.290
	189	2.4	0.10
	190	6.7	0.29
	191	6.7	0.57
20	192	0.84	0.035
	193	2	0.072
	194	2.4	0.083
	195	2.0	0.042
25	196	1.7	0.027
	197	1.2	0.16
	198	3.4	0.062
•	199	1.4	0.036
	200	1.4	0.041
30	201	1.2	0.44
	202	1.4	0.76
	203	1.0	0.036
	204	0.71	0.031
	205	11	0.036
35	206	1.7	0.095
	207	1.2	0.50
	208	2.8	0.17
	209	1.2	0.136
40	210	0.84	0.041
₩	211	0.35	0.024
	212	0.50	0.036
	213	1.0	0.55
	214	0.71	0.024
45	215	2.8	0.25
	216	0.35	0.032
	217	13	0.57
	218	1.0	0.11
	219	0.71	0.044
50	220	0.71	0.05
	221	0.71	0.041
	222	0.84	0.072
	223	0.79	0.055
EE	224	0.63	0.055
55	225	0.63	0.072

TABLE 5

5	Compound No.	Vancomycin Resistant Strains	Vancomycin Sensitive Strains
	226	1.6	0.041
10	227	0.71	0.11
70	228	1.0	0.14
	229	0.50	0.024
	230	0.35	0.031
	231	1.7	0.11
15	232	0.71	0.29
	233	1.7	1.7
	234	2	2
	235	2.4	0.25
	236	1.4	0.5
20	237	1.0	0.048
	238	1.4	0.14
	239	2.8	0.095
	240	1.19	0.055
25	241	1.4	0.048

A number of the lactic acid bacteria including all Leuconostocs, all Pediococci, and some Lactobacilli, are intrinsically resistant to vancomycin. With the increased use of vancomycin, infections due to these bacteria have been reported with increasing frequency in immunocompromised patients (Handwerger et al., Reviews of Infectious Disease 12:602-610 (1990); Ruoff et al., Journal of Clinical Microbiology 26:2064-2068 (1988)). One important aspect of the antimicrobial activity of the formula I compounds is their activity against the vancomycin-resistant lactic acid bacteria. The compounds of the present are useful in inhibiting the growth of vancomycin-resistant lactic bacteria such as Leuconostoc, Pedicocci, and Lactobacilli and thus, controlling opportunistic infections by this group of bacteria. This activity is illustrated in Table 6, which summarizes a comparison of the activity of illustrative compounds against representative vancomycin-resistant lactic acid bacteria (Pedicoccus acidilacti Pedicoccus pentosaceus, Leuconostoc lactis, Leuconostoc mesenteroides, Leuconostoc pseudomesenteroides, Leuconostoc citreum, and Lactobacillus confusus, mean geometric MIC (mcg/mL)), as determined using a standard agar dilution assay on brain-heart infusion agar.

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/ancomycin A82846B 108 51 52 Pediococcus (mean of 10) acidilacti 58 Pediococcus pentosaceus (mean of 2) Leuconostoc (mean of 2) lactis 91 64 16 mesenteroides Leuconostoc (mean of >256 76 64 2 91 32 16 11 œ ۵ pseudomesent-Leuconostoc eroides >128 >256 >128 >128 128 128 128 64 32 16 3 16 16 32 32 Leuconostoc citreum >256 >128 >128 >128 >64 128 128 128 64 32 64 32 6 6 164 Lactobacillus confusus >256 64 32 32 32 32 32 32 32 32 32 16 64 64

Table 6
In Vitro Activity of Formula I Compounds
MIC (mcg/ml)/Compound

Pharmaceutical formulations of the formula I compounds are also part of this invention. Thus, the compound, preferably in the form of a pharmaceutically acceptable salt, can be formulated for oral or parenteral administration for the therapilities or prophylactic treatment of bacterial infections.

For example, the compound can be admixed with conventional pharmaceutical carriers and excipi nts and used in the form of table ts, capsules, elixirs, suspensions, syrups, wafers, and the like. The compositions comprising a formula I compound will contain from about 0.1 to about 90% by weight of the active compound, and

mor g n rally from about 10 to about 30%. The compositions may contain common carri rs and xcipients, such as corn starch or gelatin, lactose, sucros, microcrystallin cellulose, kaolin, mannitol, dicalcium phosphate, sodium chloride, and alginic acid.

Disintegrators commonly used in the formulations of this invintion include croscarm llos , microcrystal-line cellulos , c rn starch, sodium starch glycolate and alginic acid.

Tablet binders that can be included are acacia, methylcellulose, sodium carboxymethylcellulose, polyvinylpyrrolidone (Povidone), hydroxypropyl methylcellulose, sucrose, starch and ethylcellulose.

Lubricants that can be used include magnesium stearate or other metallic stearates, stearic acid, silicone fluid, talc, waxes, oils and colloidal silica.

Flavoring agents such as peppermint, oil of wintergreen, cherry flavoring or the like can also be used.

It may be desirable to add a coloring agent to make the dosage form more attractive in appearance or to help identify the product.

For intravenous (IV) use, a water soluble form of the antibiotic can be dissolved in one of the commonly used intravenous fluids and administered by infusion. Such fluids as, for example, physiological saline, Ringer's solution, or 5% dextrose solution can be used.

For intramuscular preparations, a sterile formulation of a suitable soluble salt form of the compound, for example the hydrochloride salt, can be dissolved and administered in a pharmaceutical diluent such as pyrogen-free water (distilled), physiological saline or 5% glucose solution. A suitable insoluble form of the compound may be prepared and administered as a suspension in an aqueous base or a pharmaceutically acceptable oil base, for example, an ester of a long chain fatty acid such as ethyl oleate.

For oral use, a sterile formulation of a suitable salt form of the antibiotic, for example, the hydrochloride salt, formulated in a diluent such as distilled or deionized water, is particularly useful.

Alternatively, the unit dosage form of the antibiotic can be a solution of the antibiotic, preferably in its salt form, in a suitable diluent in sterile, hermetically sealed ampoules. The concentration of the antibiotic in the unit dosage may vary, for example, from about 1 percent to about 50 percent depending on the particular form of the antibiotic and its solubility and the dose desired by the physician.

In a further aspect, this invention provides a method for treating infectious diseases, especially those caused by Gram-positive microorganisms, in animals. The compounds of this invention are particularly useful in treating infections caused by methicillin-resistant staphylococci. Also, the compounds are useful in treating infection due to enterococci. Examples of such diseases are severe staphylococcal infections, for example, staphylococcal endocarditis and staphylococcal septicemia. The animal may be either susceptible to, or infected with, the microorganism. The method comprises administering to the animal an amount of a formula I compound which is effective for this purpose. In general, an effective amount of a formula I compound is a dose between about 0.5 and about 100 mg/kg. A preferred dose is from about 1 to about 60 mg/kg of active compound. A typical daily dose for an adult human is from about 50 mg to about 5 g.

In practicing this method, the antibiotic can be administered in a single daily dose or in multiple doses per day. The treatment regimen may require administration over extended periods of time, for example, for several days or for from one to six weeks. The amount per administered dose or the total amount administered will depend on such factors as the nature and severity of the infection, the age and general health of the patient, the tolerance of the patient to the antibiotic and the microorganism or microorganisms involved in the infection.

A convenient method of practicing the treatment method is to administer the antibiotic via intravenous infusion. In this procedure a sterile formulation of a suitable soluble salt of the antibiotic is incorporated in a physiological fluid, such as 5% dextrose solution, and the resulting solution is infused slowly IV. Alternatively, the piggy-back method of IV infusion can also be used.

In order to illustrate more fully the operation of this invention, the following examples are provided, but are not to be construed as a limitation on the scope of the invention.

EXAMPLE 1

METHOD A

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Preparation of Compound 2

A mixture of A82846B-triacetate, (2.25 g, 1.27 mmol, 1.0 equivalents (eq)) in 1:1 DMF/methanol (140 mL) under an atmosphere of argon was tr ated with 4-biphenylcarboxald hyde (331 mg, 2.12 mmol, 1.7 eq). Th resulting mixtur was heated to 70°C and maintained as such for 1.75-2 hours. Th solution was then tr ated with sodium cyanoborohydride (554 mg, 8.83 mmol, 6.9 eq). Heating at 70°C was continued for an additional 1.75-2 hours aft r which th reaction mixtur was cooled to room temperatur, conc ntrat d *in vacuo*, diluted

with water (150 mL), and lyophilized to giv a solid.

The solid was purified by preparativ revers -phas high performance liquid chromatography (HPLC) using a Waters 3 x (40 x 100 mm) C18 Nova-Pak cartridge with Waters C18 Nova-pak guard insert and utilizing TEAP buff r system. The analytical method for analysis was: 0.2% TEA/phosph ric acid (TEAP), pH = 3, th gradient syst m at tim 0 was 5% CH₃CN/94.8% H₂O with 0.2% TEAP h Id constant and at 20 minutes was 60% CH₃ON/39.8% H₂O with 0.2% TEAP held constant. The UV wavelength used was 235 nm and the flow rate was 2 ml/minute. Analysis was done using a Waters Nova-pak C18 RCM column (8 X 100mm) with a Nova-pak C18 guard insert. It is necessary to desalt the product after reverse phase purification when this HPLC method is used.

Desalting was accomplished by adding the purified product to 5-10 ml of H₂O. 1 N HCl was added dropwise with stirring to dissolve the sample. The pH at this point was approximately 1-3. The pH of the solution was then raised to 8.2 with 1 N NaOH. A white solid precipitated out of solution. The mixture was cooled, filtered, and dried under vacuum at room temperature for 8-15 hours to give the zwitter ion (or neutral compound) of the desired product, compound 2 (p-phenylbenzyl-A82846B), (1.02 g, 45%).

EXAMPLE 2

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Preparation of Compound 4

A mixture of A82846B.triacetate (1.5 g, 0.848 mmol, 1.0 eq) in methanol (100 mL) under an atmosphere of argon was treated with ρ -phenoxybenzaldehyde (298 mg, 1.51 mmol, 1.8 eq). The resulting mixture was heated to reflux and maintained as such for 2 hours. The solution was then treated with sodium cyanoborohydride (326 mg, 5.18 mmol, 6.1 eq). Heating at reflux was continued for an additional 2 hours after which the reaction mixture was cooled to room temperature and evaporated to dryness *in vacuo*.

The product was purified by reverse-phase HPLC with a TFA buffer. The analytical method for analysis was accomplished by using a Waters Nova-pak C18 RCM column (8 x 100 mm) with a Nova-pak C18 guard insert, eluting with a 2.0 ml/minute linear gradient of 15% acetonitrile/0.1% TFA at time zero to 80% acetonitrile/0.1% TFA at 15 minutes. The fractions containing the products were detected by ultraviolet scan at 235 nm. The organic solvent of the desired fractions was removed and the mixture was lyophilized to a white solid to give 0.618 mg of *p*-phenoxybenzyl-A82846B compound 4-tris(trifluroacetate) salt (20% yield). No desalting or further purification was necessary. This method is also especially useful in the preparation of Compound 2 wherein phenylbenzaldehyde is one of the starting materials.

EXAMPLE 3

Method B

Preparation of Compound 176

A mixture of A82846B.triacetate (280 mg, 0.157 mmol, 1.0 eq) in 1:1 DMF/methanol (30 mL) was treated with 8-phenyloctanal (59 mg, 0.29 mmol, 1.8 eq) and sodium cyanoborohydride (60 mg, 0.95 mmol, 6.1 eq). The resulting mixture was heated, under an atmosphere of nitrogen, to 70°C and maintained as such for 1 hour. The reaction mixture was then cooled to room temperature and concentrated in vacuo to give a residue. Purification of the product was accomplished by reverse-phase preparative HPLC utilizing a Waters 2 x (40 x 100 mm) C18 Nova-Pak cartridge with Waters C18 Nova-Pak guard insert. Elution was accomplished with a 30 minute linear gradient (time=0 minutes 95% TEAP (0.5% aqueous triethylamine adjusted to pH=3 with phosphoric acid)/5% CH₃CN to t = 30 minutes 20% TEAP/80% CH₃CN) with a flow rate of 40 mL/minute and UV detection at 280 nm. The desired fraction was concentrated in vacuo then desalted with a Waters Sep-Pak cartridge as described below. This afforded compound 176 in 22% yield (60 mg).

The resulting compound was desalted as follows. A Waters Sep-Pak cartridge was pre-wet with methanol (2-3 column volumes) then conditioned with water (2-3 column volumes). The sample, dissolved in a minimum volume of water, was loaded onto the Sep-Pak column which was then washed with water (2-3 column volumes) to remove the unwanted salts. The product was then eluted with an appropriate solvent system, typically 1:1 CH₃CN/H₂O, CH₃CN, and/or methanol. The organic solvent component was removed *in vacuo* and the resulting aqueous solution lyophilized to give the final product.

EXAMPLE 4

Preparation of Compound 229

A three lit r 3-neck d flask was fitt d with a condens r, nitrog n inl t and ov rh ad mechanical stirring apparatus. The flask was charged with pulverized A82846B acetate salt (20.0 g, 1.21 x 10⁻³ mol) and methanol (1000 mL) under a nitrogen atmosphere. 4'-chlorobiphenylcarboxaldehyde (2.88 g, 1.33 x 10⁻² mol, 1.1 eq.) was added to this stirred mixture, followed by methanol (500 mL). Finally, sodium cyanoborohydride (0.84 g, 1.33 x 10⁻² mol, 1.1 eq.) was added followed by methanol (500 mL). The resulting mixture was heated to reflux (about 65°C).

After 1 hour at reflux, the reaction mixture attained homogeneity. After 25 hours at reflux, the heat source was removed and the clear reaction mixture was measured with a pH meter (6.97 at 58.0°C). 1 N NaOH (22.8 mL) was added dropwise to adjust the pH to 9.0 (at 54.7°C). The flask was equipped with a distillation head and the mixture was concentrated under partial vacuum to a weight of 322.3 grams while maintaining the pot temperature between 40-45°C.

The distillation head was replaced with an addition funnel containing 500 mL of isopropanol (IPA). The IPA was added dropwise to the room temperature solution over 1 hour. After approximately 1/3 of the IPA was added, a granular precipitate formed. The remaining IPA was added at a faster rate after precipitation had commenced. The flask was weighed and found to hold 714.4 grams of the IPA/methanol slurry.

The flask was re-equipped with a still-head and distilled under partial vacuum to remove the remaining methanol. The resulting slurry (377.8 g) was allowed to chill in the freezer overnight. The crude product was filtered through a polypropylene pad and rinsed twice with 25 mL of cold IPA. After pulling dry on the funnel for 5 minutes, the material was placed in the vacuum oven to dry at 40°C. A light pink solid (22.87 g (theory = 22.43 g)) was recovered. HPLC analysis versus a standard indicated 68.0% weight percent of Compound 229 (4-[4-chlorophenyl]benzyl-A82846B] in the crude solid, which translated into a corrected crude yield of 69.3%.

The products of the reaction were analyzed by reverse-phase HPLC utilizing a Zorbax SB-C18 column with ultraviolet light (UV; 230 nm) detection. A 20 minute gradient solvent system consisting of 95% aqueous buffer/5% CH₃CN at time=0 minutes to 40% aqueous buffer/60% CH₃CN at time=20 minutes was used, where the aqueous buffer was TEAP (5 ml CH₃CN, 3 ml phosphoric acid in 1000 ml water).

EXAMPLE 5

Table 7 summarizes the preparation and certain physical characteristics of the exemplified compounds. The yield of the product was calculated using the amount of the formula II compound as the limiting reagent. The following terms are found in Table 6 and are defined here. "Method" refers to the method of synthesis as described in Examples 1 and 2, or 3. "Reagent Equivalents" refers to the molar equivalents of the aldehyde and reducing agent relative to the formula II compound. "FAB-MS (M+3H)" refers to Fast atom bombardment-mass spectrometry.

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TABLE 7

5	İ			Reagent Equivalents	PAB-MS
	Compound	,	Method/	•	(M+3H)
	No.	(%)	DMF: MeOH	(aldehyde/	(M+3E)
		i	•	NaBH3CN)	!
10	1	28	A/1:1	1.7/6.9	1733*
10	2	45	A/1:1	1.7/6.9	1760
	3	28	A/1:1	1.8/7.6	1732**
	4	20	A/0:1	1.8/6.1	1776***
	5	30	A/0:1	1.8/6.1	1790
15	6	10	A/0:1	1.8/6.1	1768***
	7	55	A/0:1	1.8/6.1	1740***
	8	16	A/0:1	1.8/6.1	1826
	9	32	A/0:1	1.8/6.1	1764***
	10	6	A/0:1	1.8/6.1	1868
20		38	A/0:1	1.8/6.1	1784
	12	46	A/0:1	1.8/6.1	1940
	13	32	A/0:1	1.8/6.1	1783**
	14	5.4	A/1:1	1.9/4.2	1859
	15	42	A/0:1	1.8/6.1	1763
25	16	39	A/0:1	1.8/6.1	1807**
	17	41	A/0:1	1.8/6.1	1798
	18	27	A/0:1	1.8/6.1	1817
	19	30	A/0:1	1.8/6.1	1739
30	20	5	A/1:1	1.8/1.8	1775*
00	21	11	A/1:1	1.8/1.8	1872*
	22	8	A/1:1	1.8/1.8	1829**
	23	ND	A/0:1	1.8/3.6	1888***
	24	34	A/0:1	1.7/2.5	1685
35	25	31	A/0:1	1.8/1.6	1779
	26	30	A/0:1	1.7/2.5	1685
	27	19	A/0:1	1.8/2.5	1734**
	28	35	A/0:1	1.6/1.6	1735
	29	39	A/0:1	1.6/1.6	1785**
40	30	29	A/0:1	1.6/1.6	1734**
	31	11	A/0:1	1.7/2.5	1684**
	32	28	A/0:1	1.5/1.6	1771**
	33	ND	A/1:1	1.8/1.8	1789
4E	34	ND	A/1:1	1.8/1.8	1836
45	35	ND	A/1:1	1.8/1.8	1785
	36	ND	A/1:1	1.8/1.8	1835
	37	31	A/0:1	1.5/1.5	1752***
	38	16	A/0:1	1.5/1.6	1709
50	39	46	A/0:1	1.5/1.5	1773
	40	29	A/1:1	1.8/1.8	1846*
	41	46	A/0:1	1.5/1.5	1729
	42	53	A/0:1	1.5/1.5	1780
	43	22	A/0:1	1.1.1.5	1799***
55	44	42	A/0:1	1.5/1.5	1749

TABLE 7

5	Compound	Yield	Method/	Reagent Equivalents	Pab-MS
	No.	(%)	DMF: MeOH	(aldebyde/	(M+3H)
ì				Nabh3cw)	
	45	50	A/0:1	1.1/1.5	1841
10	46	38	A/0:1	1.1/1.5	1850
	47	40	A/0:1	1.5/1.5	1687
	48	22	A/0:1	1.5/1.5	1728***
	49	44	A/0:1	1.5/1.5	1776***
45	50	32	A/1:10	2.0/1.5	1774
15	51	32	A/0:1	1.5/1.5	1820
	52	31	A/0:1	1.5/1.5	1819**
	53	43	A/0:1	1.5/1.5	1896
	54	4	A/1:1	1.8/1.8	1789
20	55	21	A/0:1	1.5/1.5	1767
	56	20	A/0:1	1.1/1.5	1741
	57	29	A/0:1	1.5/1.5	1820**
	58	22	A/0:1	1.5/1.5	1727
	59	ND	A/1:1	1.8/1.8	1803
25		33	A/0:1	1.1/1.5	1777**
	60	24	A/0:1	1.1/1.5	1723
	61	ND	A/1:1	1.8/1.8	1789**
•	62 63	ND	A/1:1	1.8/1.8	1789**
	64	30	A/0:1	1.5/1.5	1805
30	65	24	A/0:1	1.1/1.5	1763
	66	17	A/0:1	1.1/1.5	1704***
•	67	22	A/0:1	1.1/1.5	1766***
	68	ND	A/1:1	1.8/1.8	1802
35	69	ND	A/1:1	1.8/1.8	1803
	70	44	A/0:1	1.1/1.5	1821
	71	4	A/0:1	1.1/1.5	1796***
	72	32	A/0:1	1.5/1.5	1750***
	73	ND	A/1:1	1.8/1.8	1753
40	74	17	A/0:1	1.1/1.5	1815
	75	23	A/0:1	1.5/1.5	1806***
	76	16	A/1:1	1.8/1.8	1711
	77	ND	A/1:1	1.8/1.8	1742
	78	5	A/1:1	1.8/1.8	1728
45	79	ND	A/1:1	1.8/1.8	1783**
	80	46	A/0:1	1.5/1.5	1843****
	81	52	A/0:1	1.5/1.5	1844***
!	82	29	A/0:1	1.5/1.5	1726***
50	83	7	A/0:1	1.5/1.5	1798**
==	84	8	A/0:1	1.5/1.5	1700
	85	30	A/0:1	1.5/1.5	1775
	86	45	A/0:1	1.5/1.5	1809
	87	42	A/0:1	1.1/1.5	1854**
55	88	36	A/0:1	1.1/1.5	1854**

TABLE 7

		:		Reagent	
5	Compound	Yield	Method/	Equivalents	Pab-MS
	No.	(%)	DMF: MeOH	(aldehyde/	(M+3H)
	1			NaBH3CN)	ı
	89	43	A/1:1	1.8/1.8	1711
10	90	13	A/1:1	1.8/1.8	1787
	91	20	A/1:10	1.5/1.5	1759**
	92	23	A/1:10	1,5/1,5	1777
	93	42	A/0:1	1.5/1.5	1823
	94	41	A/0:1	1.1/1.5	1854**
15	95	49	A/0:1	1.1/1.5	1789**
	96	34	A/0:1	1.1/1.5	1832
	97	42	A/1:10	1.5/1.5	1773**
				1/1.5	1805
20	98	31	A/0:1		1770**
	99	ND	A/1:1 A/1:1	1.8/1.8	1787
	100	ND		1.19/1.8	1761
	101	34	A/1:1	1.5/1.5	1805
	102	41	A/0:1	1/1.5	1788***
25	103	37	A/0:1	1.1/1.5	1819**
	104	34	A/0:1	1.7/2.0	1838*
	105	ND	A/1:1		1844
	106	ND	A/1:1	1.7/2.0	1802
	107	ND	A/1:1	1.8/1.8	1791**
30	108	ND	A/0:1	1.8/1.8	1789
	109	ND	A/0:1	1.1/1.5	1881
	110	15	A/0:1	1.8/1.8	1843
	111	ND	A/1:1	1.8/1.8	1764
35	112	16	A/1:1	1.1/1.5	1805**
	113	45	A/0:1	1.1/1.5	1888**
	114	52	A/0:1		1791
	115	39	A/0:1	1.1/1.5	1834
	116	ND 29	A/1:1 A/0:1	1.5/1.7	1803**
40	117	28	A/0:1	2/1.5	1765**
	118		A/0:1	1/1.5	1843
	119	38	A/0:1	1.1/1.5	1757
	120			1.1/1.5	1799
	121	41	A/0:1		
45	122	24	A/1:1	1.8/2.6	1863
	123	55	A/0:1	1.1/1.5	1781**
	124	17	A/1:10	3/1.5	1841
	125	36	A/0:1	1.5/1.8	1818
50	126	26	A/0:1		1810
50	127	54	A/0:1	1.1/1.5	
	128	. 34	A/0:1	1.4/1.8	1831
	129	ND	A/1:1	1.4/1.8	1780
	130	4	A/0:1	1.1/1.5	1795**
55	131	42	A/0:1	1.1/1.5	1834**

TABLE 7

			Reagent	!
Compound	i	Method/	Equivalents	PAB-M
No.	(%)	DMF: MeOH:	(aldebyde/	(M+3H)
			NaBH3CN)	
132	49	A/0:1	1.1/1.5	1843
133	41	A/0:1	1.1/1.5	1855
134	30	A/0:1	1.1/1.5	1801**
135	ND	A/1:1	1.8/1.8	1779
136	ND	A/1:1	1.8/1.8	1699
137	ND	A/1:1	1.8/1.8	1760
138	ND	A/1:1	1.8/1.8	1741
139	13	A/1:10	2.4/1.5	1749**
140	11	A/1:10	2.9/1.5	1750*
141	ИD	A/1:1	2.3/5.3	1742
142	ND	A/1:1	2.5/5.4	1826
143	ND	A/1:1	1.8/1.8	1861
144	ND	A/1:1	1.5/1.5	1922
145	ND	A/1:1	1.1/1.1	1716
146	ND	A/1:1	1.35/1.8	1780*
147	ND	A/1:1	1.5/1.8	1769
148	31	A/1:10	3/1.5	1857
149	18	A/0:1	1.1/1.5	1777
150	22	A/1:1	2/4.8	1803
151	ND	A/1:1	1.8/1.8	1760
152	ND	A/1:1	1.8/1.8	1826***
153	22	A/1:10	2.5/1.6	1782
154	ND	A/1:1	1.8/1.8	1780
155	13	A/0:1	1.6/1.6	1768
156	41	A/1:9	1.2/1.6	1788
157	9	A/1:1	2.7/5.4	1810
158	ND	A/1:1	1.8/4.1	1854
159	13	A/1:9	1/1.6	1807
160	13	A/1:9	0.95/1.6	1774
161	ND	A/1:1	1.8/1.8	1690
162	ND	A/1:1	3.1/6.9	1804
163	ND	A/1:1	1.9/5.3	1854
164	ND	A/1:1	1.8/1.8	1772
165	21	A/1:1	2.0/4.9	1810
166	20	A/1:1	2.0/6.2	1870
167	23	A/1:1	1.8/4.1	1914
168	ND	A/1:1	1.8/1.8	1737
169	15	A/1:1	1.8/4.1	1700
170	39	A/0:1	1.2/1.1	1728
171	32	A/0:1	1.2/1.5	1729**
172	11	B/1:1	2.2/4.8	1755**
173	51	A/1:9	1.3/1.7	1909
174	35	A/1:9	1.5/1.6	1816

TABLE 7

				Reagent	:
5	Compound	Yield	Method/	Equivalents	Pab-Ms
	No.	(%)	DMF: MeOH	(aldebyde/	(M+3H)
				NaBH3CN)	1
	175	22	B/1:1	1.9/6.2	1742
10	176	21	B/1:1	1.8/6.1	1782
,,	177	ND	A/1:1	3.6/1.8	1774
	178	33	A/1:9	1.4/1.7	1788**
	179	22	B/1:1	1.8/3.8	1748
	180	16	A/1:1	1.1/1.3	1591***
15	181	14	A/1:1	1.1/1.3	1617
	182	17	A/0:1	1.6/6.3	1725
	183	17	A/0:1	1.6/6.3	: 1691**
	184	8	A/0:1	1.6/6.26	1707**
	185	21	A/1:1	1.1/3.0	1725**
20	186	8	A/1:1	1.1/3.0	1630**
	187	16	A/1.1	1.6/3.0	2110**
	188	6	A/1.1	1.5/5.0	2976**
	189	20	A/1:10	1/1.2	1747**
05	190	9	A/1:10	1.5/1.5	1716
25	191	18	B/1:1	1.8/4.1	1771**
	192	11	A/0:1	ND/1.8	1738
	193	24	A/1:10	2.0/1.5	1820**
	194	27	A/1:10	2.0/1.5	1821
30	195	18	B/1:1	1.6/3.6	1798
	196	18	B/1:1	1.8/3.9	1754
	197	35	B/1:1	1.5/3.5	1810
	198	14	B/1:1	1.5/3.7	1784
	199	ND	B/1:1	1.5/2.8	1772
35	200	11	B/1:1	1.5/3.7	1828
	201	14	B/1:1	1.8/6.3	1873**
	202	7	B/1:1	1.3/5.9	1889**
	203	15	A/0:1	1.1/1.1	1843
40	204	16	B/1:1	2.0/5.6	1746
40	205	23	B/1:1	1.8/3.7	1732
	206	11	A/0:1	1.1/1.1	1777
	207	11	B/1:1	1.6/4.2	1813**
	208	26	B/1:1	1.9/3.9	1703
45	209	20	A/1:1	1.0/1.6	1774
	210	35	A/0:1	1.0/1.0	1788
	211	26	A/0:1	1.3/1.8	1777
	212	48	A/1:1	1.1/3.1	1849**
	213	56	A/1:1	1.0/3.6	1849**
50	214	9	B/1:1	1.9/1.9	1732
	215	35	A/0:1	1.3/1.8	1820***
	216	31	A/0:1	1.3/1.8	1828***
	217	12	B/1:1	2.0/2.1	1676
	218	24	A/1:10	1.2/1.5	1766***
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TABLE 7

Compound No.	Yield (%)	Method/	Reagent Equivalents (aldehyde/ NaBH3CN)	PAB-MS (M+3H)
219	24	A/1:1	1.4/3.5	1860
220	21	A/0:1	1.3/1.8	1785
221	42	A/0:1	1.3/1.8	1787
222	20	A/0:1	1.1/1.1	1787
223	32	A/1:1	2.4/4.5	1817**
224	36	A/1:1	1.6/5.6	1773**
225	ND	A/0:1	1.1/1.1	1787
226	28	A/1:1	1.5/3.0	1766*
227	22	A/1:1	1.2/3.7	1777**
228	21	A/0:1	1/1.1	1848**
229	16	A/0:1	1/1.2	1793
230	27	A/0:1	1.3/1.8	1838***
231	36	A/0:1	1.3/1.8	1785*
232	32	A/1:1	1.8/4.6	1806
233	5	A/1:1	1.1/7.3	1878
234	7	B/1:1	1.5/3.5	1836*
235	15	B/1:1	1.4/4.8	1750
236	4	B/1:1	1.4/6.3	1819**
237	14	A/0:1	1.1/1.1	1787
238	25	B/0:1	1.1/1.1	1771
239	22	B/1:1	1.6/1.5	1810
240	4.7	A/1:60	1.2/1.1	1810**
241	24	B/1:1	1.1/2.5	1779**
242	N.D.	A/1:50	1.1/1.2	1787
243	20	A/0:1	1.1/1.1	1790
244	24	C/0:1	1.1/1.1	1808
N.D.= Not	determ	ined		1
*M+H				
**M+2H			_	
***M+4H				
****M+6H				

EXAMPLE 6

Capsule Formulation

Capsules containing 250 mg of Compound 2 are prepared using the following ingredients:

Ingredient	Weight
Compound 2 HCl salt	255.4 mg
Corn starch flowable powder	150 mg
Corn starch	144.6 mg

Compound 2 (HCl salt form, 255.4 mg), corn starch flowable powder (150 mg) and corn starch (144.6 mg) ar

blended in a suitable mix r until homogenous. The mixture is used to fill a hard gelatin capsule to a nit fill wight of 550 mg.

EXAMPLE 7

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Capsule Formulation

Capsules containing 250 mg of Compound 229 are prepared using the following ingredients:

Ingredient	Weight
Compound 229 HCl salt	255.4 mg
Corn starch flowable powder	150 mg
Corn starch	144.6 mg

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Compound 2 (HCl salt form, 255.4 mg), corn starch flowable powder (150 mg) and corn starch (144.6 mg) are blended in a suitable mixer until homogenous. The mixture is used to fill a hard gelatin capsule to a net fill weight of 550 mg.

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EXAMPLE 8

Suspension Formulation

A sterile insoluble form of compound 2 is milled or screened to a particle size suitable for suspension. This particulate material is suspended in the following vehicle:

Ingredient	Weight
Lecithin	1%
Sodium citrate	2%
Propylparaben	0.015%
Distilled water	q.s. to desired volume

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EXAMPLE 9

60 Suspension Formulation

A sterile insoluble form of compound 229 is milled or screened to a particle size suitable for suspension. This particulate material is suspended in the following vehicle:

Ingredient	Weight	
Lecithin	1%	
Sodium citrate	2%	
Propylparaben	0.015%	
Distilled water	q.s. to desired volume	

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EXAMPLE 10

Tabl t Formulation

5 Tablets containing 250 mg of compound 2 are prepared with the following composition:

Ingredient	Weight
Lecithin	1%
Sodium citrate	2%
Propylparaben	0.015%
Distilled water	q.s. to desired volume

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EXAMPLE 11

Tablet Formulation

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Tablets containing 250 mg of compound 229 are prepared with the following composition:

Ingredient	Weight	
Lecithin	1%	
Sodium citrate	2%	
Propylparaben	0.015%	
Distilled water	q.s. to desired volume	

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EXAMPLE 12

35 Tablet Formulation

Tablets containing 250 mg of compound 2 are prepared with the following composition:

Ingredient	Weight
Compound 2 HCl salt	255.4 mg
Microcrystalline cellulose	101.1 mg
Croscarmellose sodium	12.0 mg
Providone	12.0 mg
Magnesium stearate	3.0 mg
Stearic acid	4.0 mg
Purified water	0.16 ml

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EXAMPLE 13

Tablet Formulation

Tablets containing 250 mg of compound 229 are prepared with the following composition:

Ingredient	W ight
Compound 229 HCI salt	255.4 mg
Microcrystalline cellulose	101.1 mg
Croscarmellose sodium	12.0 mg
Providone	12.0 mg
Magnesium stearate	3.0 mg
Stearic acid	4.0 mg
Purified water	0.16 ml

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Claims

A compound of the formula:

25 СН2ОН 30 35 40 OR⁵

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or salt thereof, wherein:

X and Y are each independently hydrogen or chloro;

R is hydrogen, 4-epi-vancosaminyl, actinosaminyl, or ristosaminyl;

R1 is hydrogen, or mannose;

R² is -NH₂, -NHCH₃, or-N(CH₃)₂;

R3 is -CH₂CH(CH₃)₂, [p-OH, m-Cl]phenyl, p-rhamnose-phenyl, [p-rhamnose-galactose]phenyl, [pgalactose-galactose]phenyl, or [p-CH3O-rhamnose]phenyl;

R4 is -CH₂(CO)NH₂, benzyl, [p-OH]phenyl, or [p-OH, m-Cl]phenyl;

R⁵ is hydrogen, or mannose;

R⁶ is 4-epi-vancosaminyl, L-acosaminyl, L-ristosaminyl, or L-actinosaminyl;

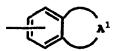
 $\mathsf{R}^{7}\,\mathsf{is}\,(\mathsf{C}_{2}-\mathsf{C}_{18})\mathsf{alkenyl},\,(\mathsf{C}_{2}-\mathsf{C}_{12})\mathsf{alkynyl},\,(\mathsf{C}_{1}-\mathsf{C}_{12}\,\mathsf{alkyl})-\mathsf{R}_{8},\,(\mathsf{C}_{1}-\mathsf{C}_{12}\,\mathsf{alkyl})-\mathsf{halo},\,(\mathsf{C}_{2}-\mathsf{C}_{6}\,\mathsf{alkenyl})-\mathsf{R}_{8},\,(\mathsf{C}_{2}-\mathsf{C}_{18})-\mathsf{halo},\,(\mathsf{C}_{2}-\mathsf{C}_$ C₆ alkynyl)-R₈, (C₁-C₁₂ alkyl)-O-R₈, and is attached to the amino group of R⁶;

R8 is s lected from the group consisting of:

a) multicyclic aryl unsubstitut d or substituted with on or mor substitu nts ind pendently select d from the group consisting of:

(i) hydroxy,

- (ii) halo,
- (iii) nitro,
- (iv) (C₁-C₆)alkyl,
- (v) (C₁-C₆)alkenyl,
- (vi) (C₁-C₆)alkynyl,
- (vii) (C1-C6)alkoxy,
- (viii) halo-(C₁-C₆)alkyl,
- (ix) halo-(C1-C6)alkoxy,
- (x) carbo-(C₁-C₆)alkoxy,
- (xi) carbobenzyloxy,
- (xii) carbobenzyloxy substituted with (C_1-C_8) alkyl, (C_1-C_8) alkoxy, halo, or nitro,
- (xiii) a group of the formula -S(O)_n-R⁹, wherein n' is 0-2 and R⁹ is (C₁-C₆)alkyl, phenyl, or phenyl substituted with (C₁-C₆)alkyl, (C₁-C₆)alkoxy, halo, or nitro, and
- (xiv) a group of the formula -C(O)N(R10)2 wherein each R10 substituent is independently hydrogen,
- (C_1-C_8) -alkyl, (C_1-C_6) -alkoxy, phenyl, or phenyl substituted with (C_1-C_6) -alkyl, (C_1-C_6) -alkoxy, halo,
- b) heteroaryl unsubstituted or substituted with one or more substituents independently selected from the group consisting of:
 - (i) halo,
 - (ii) (C₁-C₆)alkyl,
 - (iii) (C1-C8)alkoxy,
 - (iv) halo-(C1-C6)alkyl,
 - (v) halo-(C₁-C₆)alkoxy,
 - (vi) phenyl,
- (vii) thiophenyl,
 - (viii) phenyl substituted with halo, (C₁-C₆)alkyl, (C₁-C₆)alkenyl, (C₁-C₆)alkynyl, (C₁-C₆)alkoxy, or nitro,
 - (ix) carbo-(C₁-C₆)alkoxy,
 - (x) carbobenzyloxy,
 - (xi) carbobenzyloxy substituted with (C₁-C₆)alkyl, (C₁-C₆) alkoxyl, halo, or nitro,
 - (xii) a group of the formula -S(O)n-R9, as defined above,
 - (xiii) a group of the formula -C(O)N(R10)2 as defined above, and
 - (xiv) thienyl;
 - c) a group of the formula:



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wherein A^1 is $-OC(A^2)_2-C(A^2)_2-O$, $-O-C(A^2)_2-O$, $-C(A^2)_2-O$, or $-C(A^2)_2-C(A^2)_2-C(A^2)_2-C(A^2)_2$, and each A2 substituent is independently selected from hydrogen, (C1-C6)-alkyl, (C1-C6)alkoxy, and (C₄-C₁₀)cycloalkyl;

d) a group of the formula:

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wherein p is from 1 to 5; and

R¹¹ is independently selected from the group consisting of:

- (i) hydrogen,
- (ii) nitro.
- (iii) hydroxy,
- (iv) halo,
- (v) (C₁-C₈)alkyl,

(vi) (C₁-C₈)alkoxy,

(vii) (C9-C12)alkyl,

(viii) (C2-C9)alkynyl,

(ix) (C₉-C₁₂)alkoxy,

(x) (C_1-C_3) alkoxy substitut d with (C_1-C_3) alkoxy, hydroxy, halo (C_1-C_3) alkoxy, or (C_1-C_4) alkylthio,

(xi) (C₂-C_δ)alkenyloxy,

(xii) (C₁-C₁₃)alkynyloxy

(xiii) halo-(C₁-C₆)alkyl,

(xiv) halo-(C₁-C₆)alkoxy,

(xv) (C2-C6)alkylthio,

(xvi) (C2-C10)alkanoyloxy,

(xvii) carboxy-(C2-C4)alkenyl,

(xviii) (C₁-C₃)alkylsulfonyloxy,

(xix) carboxy-(C1-C3)alkyl,

(xx) N-[di(C_1 - C_3)-alkyl]amino-(C_1 - C_3)alkoxy,

(xxi) cyano-(C1-C6)alkoxy, and

(xxii) diphenyl-(C₁-C₆)alkyl,

with the proviso that when R^{11} is (C_1-C_8) alkyl, (C_1-C_8) alkoxy, or halo, p must be greater or equal to 2, or when R^7 is (C_1-C_3) alkyl)- R^8 then R^{11} is not hydrogen, (C_1-C_8) alkyl, (C_1-C_8) alkoxy, or halo;

e) a group of the formula:

(R¹²)_q
(Z-R¹³)_z

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wherein q is 0 to 4;

R12 is independently selected from the group consisting of:

(i) halo,

(ii) nitro,

(iii) (C1-C6)alkyl,

(iv) (C1-C6)alkoxy,

(v) halo-(C₁-C₆)alkyl,

(vi) halo-(C1-C6)alkoxy, and

(vii) hydroxy, and

(vii) (C₁-C₆)thioalkyl;

r is 1 to 5; provided that the sum of q and r is no greater than 5;

Z is selected from the group consisting of:

(i) a single bond,

(ii) divalent (C₁-C₆)alkyl unsubstituted or substituted with hydroxy, (C₁-C₆)alkyl, or (C₁-C₆)alkoxy,

(iii) divalent (C2-C6)alkenyl,

(iv) divalent (C2-C6)alkynyl, or

(v) a group of the formula -($C(R^{14})_2$)s- R^{15} - or - R^{15} -($C(R^{14})_2$)s-, wherein s is 0-6; wherein each R^{14} substituent is independently selected from hydrogen, (C_1 - C_6)-alkyl, or (C_4 - C_{10}) cycloalkyl; and R^{15} is selected from -O-, -S-, -SO₂-, -SO₂-O-, -C(O)-, -OC(O)-, -C(O)O-, -NH-, -N(C₁- C_6 alkyl)-, and -C(O)NH-, -NHC(O)-, N=N;

R¹³ is independently selected from the group consisting of:

(i) (C₄-C₁₀)heterocyclyl,

(ii) heteroaryl,

(iii) (C₄-C₁₀)cycloalkyl unsubstituted or substituted with (C₁-C₆)alkyl, or

(iv) phenyl unsubstituted or substituted with 1 to 5 substituents independently selected from: halo, hydroxy, nitro, (C_1-C_{10}) alkyl, (C_1-C_{10}) alkoxy, halo- (C_1-C_3) alkoxy, halo- (C_1-C_3) alkyl, (C_1-C_3) alkoxyphenyl, phenyl- (C_1-C_3) alkyl, (C_1-C_6) alkoxyphenyl, phenyl- (C_1-C_3) alkyl, and (C_1-C_6) alkyl-ph nyl;

f) (C_4-C_{10}) cycloalkyl unsubstituted or substituted with on or mor substituents independently selected from the group consisting of:

(i) (C₁-C₆)alkyl,

- (ii) (C₁-C₆)alkoxy,
- (iii) (C₁-C₆)alkenyl,
- (iv) (C₁-C₆)alkynyl,
- (v) (C₄-C₁₀)cycloalkyl,
- (vi) phenyl,
- (vii) phenylthio,
- (viii) phenyl substituted by nitro, halo, (C1-C8)alkanoyloxy, or carbocycloalkoxy, and
- (ix) a group represented by the formula -Z-R¹³ wherein Z and R¹³ are as defined above; and g) a group of the formula:

1 1 (R¹⁶) u

wherein

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A³ and A⁴ are each independently selected from

- (i) a bond,
- (ii) -O-,
- (iii) -S(O),-, wherein t is 0 to 2,
- (iv) -C(R^{17})₂-, wherein each R^{17} substituent is independently selected from hydrogen, (C_1 - C_6)alkyl, hydroxy, (C_1 - C_6)alkyl, (C_1 - C_6)alkoxy, or both R^{17} substituents taken together are O,
- (v) -N(R18)₂-, wherein each R18 substituent is independently selected from hydrogen; (C₁-C₆)alkyl;
- (C_1-C_6) alkenyl; (C_1-C_6) alkynyl; (C_4-C_{10}) cycloalkyl; phenyl; phenyl substituted by nitro, halo, (C_1-C_6) alkanoyloxy; or both R¹⁸ substituents taken together are (C_4-C_{10}) cycloalkyl;

R¹⁶ is R¹² or R¹³ as defined above; and u is 0-4.

2. A compound of the formula:

R⁷-R⁶-O

CH₂OH

OR

HO

NH

NH

OR

R⁴

R³

or salt thereof, wherein:

X and Y are each independently hydrogen or chloro;

R is hydrogen, 4-epi-vancosaminyl, actinosaminyl, or ristosaminyl;

R1 is hydrog n, or mannose;

 R^2 is -NH₂, -NHCH₃, or-N(CH₃)₂;

 R^3 is $-CH_2CH(CH_3)_2$, ph nyl, [p-OH,m-Cl]phenyl, p-rhamnose-phenyl, or [p-rhamnose-galactose]phenyl;

R4 is -CH₂(CO)NH₂, b nzyl, [p-OH]ph nyl, or [p-OH, m-Cl]phenyl;

R5 is hydrog n, or mannose;

R⁶ is 4-epi-vancosaminyl, L-acosaminyl, L-ristosaminyl, or L-actinosaminyl;

R7 is -(CH₂)_n-R8, or -C(CH₃)CH-R8, and is attach d to th amino group of R6;

n is 1-10;

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R⁸ is selected from the group consisting of:

- a) multicyclic aryl unsubstituted or substituted with one or more substituents independently selected from the group consisting of:
 - (i) hydroxy,
 - (ii) halo,
 - (iii) nitro,
 - (iv) (C1-C6)alkyl,
 - (v) (C₁-C₆)alkenyl,
 - (vi) (C₁-C₆)alkynyl,
 - (vii) (C1-C6)alkoxy,
 - (viii) halo-(C1-C6)alkyl,
 - (ix) halo-(C₁-C₆)alkoxy,
 - (x) carbo-(C₁-C₆)alkoxy,
 - (xi) carbobenzyloxy,
 - (xii) carbobenzyloxy substituted with (C₁-C₆)alkyl, (C₁-C₆)alkoxy, halo, or nitro,
 - (xiii) a group of the formula $-S(O)_{n'}-R^9$, wherein n' is 0-2 and R^9 is (C_1-C_6) alkyl, phenyl, or phenyl substituted with (C_1-C_6) alkyl, (C_1-C_6) alkoxy, halo, or nitro, and
 - (xiv) a group of the formula -C(O)N(R10)2 wherein each R10 substituent is independently hydrogen,
 - (C_1-C_6) -alkyl, (C_1-C_6) -alkoxy, phenyl, or phenyl substituted with (C_1-C_6) -alkyl, (C_1-C_6) -alkoxy, halo, or nitro:
- b) heteroaryl unsubstituted or substituted with one or more substituents independently selected from the group consisting of:
 - (i) halo,
 - (ii) (C1-C6)alkyl,
 - (iii) (C₁-C₆)alkoxy,
 - (iv) halo-(C1-C6)alkyl,
 - (v) halo-(C₁-C₆)alkoxy,
 - (vi) phenyl,
 - (vii) thiophenyl,
 - (viii) phenyl substituted with halo, (C_1-C_6) alkyl, (C_1-C_6) alkenyl, (C_1-C_6) alkynyl, (C_1-C_6) alkoxy, or nitro,
 - (ix) carbo-(C₁-C₆)alkoxy,
 - (x) carbobenzyloxy,
 - (xi) carbobenzyloxy substituted with (C₁-C₆)alkyl, (C₁-C₆) alkoxy, halo, or nitro,
 - (xii) a group of the formula -S(O)n-R9, as defined above, and
 - (xiii) a group of the formula -C(O)N(R10)2 as defined above;
- c) a group of the formula:

A¹

wherein A¹ is $-OC(A^2)_2-C(A^2)_2-O$, $-O-C(A^2)_2-O$, or $-C(A^2)_2-O$, or $-C(A^2)_2-C(A^2)_2-C(A^2)_2$.

and each A² substituent is independently selected from hydrogen, (C₁-C₆)-alkyl, (C₁-C₆)alkoxy, and (C₄-C₁₀)cycloalkyl;

d) a group of the formula:

wher in p is from 1 to 5; and

R¹¹ is independ ntly selected from the group consisting of:

(i) nitro,

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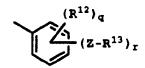
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- (ii) hydroxy,
- (iii) (C₉-C₁₂)alkyl,
- (iv) (C9-C12)alkoxy,
- (v) (C₂-C₅)alkenyloxy,
- (vi) halo-(C1-C6)alkyl,
- (vii) halo-(C₁-C₆)alkoxy,
- (viii) (C₂-C₆)alkylthio,
- (ix) (C₁-C₆)alkynyl,
- (x) (C₂-C₁₀)alkanoyloxy,
- (xi) carboxy-(C2-C4)alkenyl,
- (xii) (C₁-C₃)alkylsulfonyloxy,
- (All) (C1-C3)alkyisullollyioxy
- (xiii) carboxy-(C₁-C₃)alkyl,
- (xiv) (C₁-C₃)alkoxy substituted with (C₁-C₃)alkoxy, hydroxy, halo(C₁-C₃)alkoxy, or (C₁-C₄)alkylthio,
- (xv) N-[di(C_1 - C_3)-alkyl]amino-(C_1 - C_3)alkoxy,
- (xvi) cyano-(C₁-C₆)alkoxy,
- (xvii) (C₁-C₁₂)alkyl, (C₁-C₁₂)alkoxy, or halo when p is greater or equal to 2,
- (xviii) diphenyl-(C₁-C₆)alkyl, and
 - (xix) hydrogen, (C₁-C₆)alkyl, or (C₁-C₆)alkoxy when n greater or equal to 4;
- e) a group of the formula:



wherein q is 0 to 4;

R12 is independently selected from the group consisting of:

- (i) halo,
- (ii) nitro,
- (iii) (C1-C6)alkyl,
- (iv) (C₁-C₆)alkoxy,
- (v) halo-(C₁-C₆)alkyl,
- (vi) halo-(C₁-C₆)alkoxy, and
- (vii) hydroxy, and
- (vii) (C₁-C₆)thioalkyl;

r is 1 to 5; provided that the sum of q and r is no greater than 5;

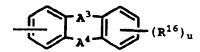
Z is selected from the group consisting of:

- (i) a single bond,
- (ii) divalent (C₁-C₆)alkyl unsubstituted or substituted with hydroxy, (C₁-C₆)alkyl, or (C₁-C₆)alkoxy,
- (iii) divalent (C2-C6)alkenyl,
- (iv) divalent (C2-C6)alkynyl, or
- (v) a group of the formula - $(C(R^{14})_2)_s$ - R^{15} or - R^{15} - $(C(R^{14})_2)_s$ -, wherein s is 0-6; each R^{14} substituent is independently selected from hydrogen, (C_1-C_6) -alkyl, or (C_4-C_{10}) cycloalkyl; and R^{15} is selected from -O-, -S-, -SO-, -SO₂-, -SO₂-O-, -C(O)-, -C(O)-, -C(O)O-, -NH-, -N(C₁-C₆ alkyl)-, and -C(O)NH-;

R¹³ is independently selected from the group consisting of:

- (i) (C₄-C₁₀)heterocyclyl,
- (ii) heteroaryl,
- (iii) (C₄-C₁₀)cycloalkyl unsubstituted or substituted with (C₁-C₆)alkyl, or
- (iv) phenyl unsubstituted or substituted with 1 to 5 substituents independently selected from: halo, hydroxy, nitro, (C_1-C_{10}) alkyl, (C_1-C_{10}) alkoxy, halo- (C_1-C_3) alkoxy, halo- (C_1-C_3) alkyl, (C_1-C_3) alkoxy-ph nyl, phenyl- (C_1-C_3) alkyl, (C_1-C_6) alkoxy-phenyl nyl- (C_1-C_3) alkyl, (C_1-C_6) alkyl-phenyl
- f) (C_4-C_{10}) cycloalkyl unsubstituted or substituted with one or mor substituents independently silected from the group consisting of:

- (i) (C₁-C₆)alkyl,
- (ii) (C₁-C₆)alk xy,
- (iii) (C1-C8)alk nyl,
- (iv) (C1-C8)alkynyl,
- (v) (C₄-C₁₀)cycl alkyl,
- (vi) phenyl,
- (vii) phenylthio,
- (viii) phenyl substituted by nitro, halo, (C₁-C₆)alkanoyloxy, or carbocycloalkoxy, and
- (ix) a group represented by the formula -Z-R13 wherein Z and R13 are as defined above; and
- g) a group of the formula:



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wherein

A³ and A⁴ are each independently selected from

- (i) a bond.
- (ii) -O-,

S(iii) -(O),-, wherein t is 0 to 2,

- (iv) -C(R^{17})₂-, wherein each R^{17} substituent is independently selected from hydrogen, (C_1 - C_6)alkyl, hydroxy, (C_1 - C_6)alkyl, (C_1 - C_6)alkoxy, or both R^{17} substituents taken together are O,
- (v) -N(R¹⁸)₂-, wherein each R¹⁸ substituent is independently selected from hydrogen; (C_1 - C_6)alkyl; (C_1 - C_6)alkenyl; (C_1 - C_6)alkynyl; (C_4 - C_1 0)cycloalkyl; phenyl; phenyl substituted by nitro, halo, (C_1 - C_6)alkanoyloxy; or both R¹⁸ substituents taken together are (C_4 - C_1 0)cycloalkyl;

 R^{16} is R^{12} or R^{13} as defined above; and u is 0-4.

- 30 3. A compound of Claim 1 wherein R is 4-epi-vancosaminyl, R¹ is hydrogen, R² is NHCH₃, R³ is CH₂CH(CH₃)₂, R⁴ is CH₂(CO)NH₂, R⁵ is hydrogen, R⁶ is 4-epi-vancosaminyl, and X and Y are Cl.
 - 4. A compound of Claim 2 wherein R is 4-epi-vancosaminyl, R¹ is hydrogen, R² is NHCH₃, R³ is CH₂CH(CH₃)₂, R⁴ is CH₂(CO)NH₂, R⁵ is hydrogen, R⁶ is 4-epi-vancosaminyl, and X and Y are Cl.

5. The compound 4-[4-chlorophenyl]benzyl-A82846B.

6. A pharmaceutical composition comprising a compound of Claim 1 to 5 or a pharmaceutically acceptable salt thereof, associated with one or more pharmaceutically acceptable carriers therefor.

7. A pharmaceutical composition as claimed in Claim 6 for use in treating susceptible bacterial infections.

- 8. A process for the preparation of a compound of any one of Claims 1 to 5 which comprises a) reacting in methanol at about 25°C to about 100°C under an inert atmosphere:
- i) a glycopeptide antibiotic of the formula:

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wherein X and Y are each independently hydrogen or chloro;

R is hydrogen, 4-epi-vancosaminyl, actinosaminyl, or ristosaminyl;

R1 is 4-epi-vancosaminyl, acosaminyl, ristosaminyl, 4-keto-vancosaminyl, or vancosaminyl;

R² is hydrogen, or mannose;

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R3 is -NH₂, -NHCH₃, or-N(CH₃)₂;

R⁴ is -CH₂CH(CH₃)₂, [p-OH,m-Cl]phenyl, p-rhamnose-phenyl, [p-rhamnose-galactose]phenyl, [p-galactose-galactose]phenyl, or [p-CH₃O-rhamnose]phenyl;

R5 is -CH2(CO)NH2, benzyl, [p-OH]phenyl, or [p-OH, m-Cl]phenyl;

R6 is hydrogen, or mannose, with

ii) an aldehyde corresponding to the group R⁷ as defined in Claim 1 at about 25°C to about 100°C; b) continuing the reaction until formation of a Schiff's base; and

c) reducing the Schiff's base by addition of a metal borohydride to the mixture at 25°C to about 100°C.

- 9. A process for the preparation of a compound of any one of Claim 1 to 5 which comprises reacting in a polar solvent at about 25°C to about 100°C under an inert atmosphere:
 - i) a glycopeptide antibiotic of the formula:

wh r in X and Y ar ach indep nd ntly hydrogen or chloro; R is hydrog n, 4-epi-vancosaminyl, actinosaminyl, or ristosaminyl; R1 is 4-epi-vancosaminyl, acosaminyl, ristosaminyl, 4-keto-vancosaminyl, or vancosaminyl; R2 is hydrog n, or mannos; R^3 is $-NH_2$, $-NHCH_3$, $r-N(CH_3)_2$; 5 R^4 is $-CH_2CH(CH_3)_2$, [p-OH,m-Cl]phenyl, p-rhamnose-phenyl, [p-rhamnose-galactose]phenyl, [p-galactose-galactose]phenyl, or [p-CH₃O-rhamnose]phenyl; R⁵ is -CH₂(CO)NH₂, benzyl, [p-OH]phenyl, or [p-OH, m-Cl]phenyl; R⁶ is hydrogen, or mannose, with ii) an aldehyde corresponding to the group R7 as defined in Claim 1, in the presence of 10 iii) a reducing agent selected from a metal borohydride, and a homogeneous or heterogeneous catalytic hydrogenation agent or agents; for a time sufficient to produce a compound of Claim 1. 10. The process of Claim 9 wherein the reducing agent is sodium cyanoborohydride, and the reaction is car-15 ried out for about 20 to 28 hours at a temperature of about 60°C to about 70°C. 11. The process of Claim 9 wherein the aldehyde is 4'biphenylcarboxaldehyde. 20 25 30 35 40 45 50



EUROPEAN SEARCH REPORT

Application Number EP 95 30 0429

Category	Citation of document with indic of relevant passa		Relevant to claim	CLASSIFICATION OF THE APPLICATION (Int.CL6)
A :	JOURNAL OF ANTIBIOTIC vol.42, no.1, January page 63-72 R NAGARAJAN ET AL. 'S antibacterial evaluat vancomycins' * the whole document	1989, TOKYO JP ynthesis and ion of N-alkyl	1-10	C07K9/00 A61K38/14
X	EP-A-0 201 251 (ELI L 1986 * the whole document	•	1-10	
D,A	EP-A-O 435 503 (ELI L * the whole document -		1-10	
				TECHNICAL FIELDS SEARCHED (int.Cl.6)
			C07K A61K	
	The present search report has been	drawn up for all claims		
	Place of search	Date of completion of the se	reb	Examiner
	THE HAGUE	9 May 1995	Mas	sturzo, P
CATEGORY OF CITED DOCUMENTS X: particularly relevant if taken alone Y: particularly relevant if combined with another document of the same category A: technological background		E : earlier pa after the D : documen	principle unduring the stant document, but publifiling date t-cited in the application t-cited for other reasons	lished on, or